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Lerman, M.I., Latif, F., Wei, M.-H., Duh, F.-M., Minna, J.D., and Gao, B.
                                                                                    Calcium channel compositions and methods Patent: US 6441156-A 37 27-AUG-2002;
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Stinchcomb, D.T., Jarvis, T. and McSwiggen, J.
Nucleic acid based inhibition of CD40
Patent: US 6194150-A 790 27-FEB-2001;
Location/Qualifiers
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; Mismatches 0;
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J. 1.1e+04;
J. 1.2e+04;
                                                                                                                                                  4.2e+04;
thes 0;
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                                               PAT 16-MAY-2001
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BD166119/c
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                                                                                                                                                              nucleic acid probes, method nucleic acid by using the probes, obtained by the method.

BD166119

BD166119
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Sequence 79
AR132367
AR132367.1
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(bases 1 to 19)

Kurane, R., Kanagawa, T., Kamagata, Y., Torimura, M., Kura Yamada, K. and Yokomaku, T.

Novel nucleic acid probes, method for determining conc nucleic acid by using the probes, and method for analy obtained by the method

Patent: JP 2002191372-A 99 09-JUL-2002;

PANTIONAL INSTITUTE OF ADVANCED INDUSTRIAL SCIENCE AND KANKYO ENGINEERING CO LTD

CS Artificial Sequence
                                                                                                                          BD166119.1 GI:27871931 JP 2002191372-A/99. unidentified unidentified
                                                                                                                                                                                                                                                                                                                                                                                                               1 (bases 1 to 15)
Stinchcomb, D.T., Jarvis, T. and McSwiggen, J.
Nucleic acid based inhibition of CD40
Patent: US 6194150-A 792 27-FEB-2001;
Location/Qualifiers
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Stinchcomb, D.T., Jarvis, T. and McSwiggen, J.
Nucleic acid based inhibition of CD40
Patent: US 6194150-A 791 27-FEB-2001;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unknown.
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Unclassified.
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Similarity 100.0%;
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/mol_type="unassigned
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Pred. No.
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Pred. No.
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           Candida albicans
Candida albicans
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Eukaryota; Fungi; mitosporic Saccharomycetales; Candida.
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AX487528
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        Roemer,T., Jiang,B.,
                                                                                                     AX487528.1 GI:22321676
                                                                                                                                               AX487528
                                                                                                                                                                                                                                                                                                                                                                                                  Patent: WO 9846645-A 19 22-OCT-1998; KUFER PETER (DE); RAUM TOBIAS (DE)
                                                                                                                                                                                                                                                                                                                                                                                                                                1 (bases 1 to 24)
Kufer, P. and Raum, T.
NOVEL METHOD FOR THE PRODUCTION OF ANTI-HUMAN ANTIGEN RECEPTORS AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19 from Patent WO9846645. A84252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       unclassified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  unidentified
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CC The sequence hybridizes with the sequence of the above no. SEH Key Location/Qualifiers
FT Source 1.19
/organism='Artificial Segmence'.
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14; Conservative
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Conservative (
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RYUICHIRO KURANE, TAKAHIRO KANAGAWA, YOICHI KAMAGATA, MASAKI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               JP 2002191372-A/99
09-JUL-2002
                                                                                                                              4828 from Patent
                                                                                                                                                                                                                                                                                                                                   /organism="unidentified"
/mol_type="unassigned DNA"
/db_xref="taxon:32644"
                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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100.0%; Pred. No. 4.:
tive 0; Mismatches
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100.0%; Pred. No. 4.2e+04;
ive 0; Mismatches 0;
       Boone, C.,
                                                                                                                               24 bp DNA
WO02053728.
       Bussey, H.
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O. 4.2e+04;
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    and Ohlsen, K.L.
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BD075290/c
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Homo sapiens
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Query Match .
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                                      T cell antigen receptor-beta chain {rearranged 8-wk fetal thymus, sample 2, mRNA, 27 nt].
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MICROMET AG
OS ATÉMICA
PN JP 2001
PD 23-OCT-
PF 14-APR-1
PR 14-APR
S81367.1 GI:245125
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  synthetic construct synthetic construct
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      artificial sequences.
1 (bases 1 to 24)
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JP 2001519824-A/19.
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BD075290
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Patent: WO 02053728-A 4828 11-JUL-2002;
Elitra Pharmaceuticals, Inc. (US)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                            GTGAGCTCAACCTG 198
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14-APR-1998 JP 1998543494
14-APR-1998 EP 97106109.8
PETER KUFER, TOBIAS RAUM
CO7K16/00,C07K16/30, A61K39/395
Description of Artificial Sequence:primer
Kev Location/Qualifiers
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JP 2001519824-A/19
23-OCT-2001
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/mol_type="genomic DNA"
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/mol_type="unassigned DNA"
/db_xref="taxon:5476"
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 14;
Pred. No.
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4.2e+04;
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                                                                                                                       PRI 07-MAY-1993
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Matsuki, Y., Horai, S. and Yabusaki, Y.
THERMOSTABLE ESTERASE
JOURNAL PATENT: JP 1995213280-A 26 15-AUG-1995;
SUMITOMO CHEM CO LTD
OS None
OC Artificial sequences.
PN JP 1995213280-A/26
PD 15-AUG-1995
PF 02-FEB-1994 JP 1994011014
PI MATSUKI YASUSHI, HORAI SHINJI, YABUSAKI YOSHIYASU PC C12N9/16, C12N1/21, C12N15/00, (C12N9/16, C12N1/21, PC C12R1:19);
CC strandedness: Single;
CC topology: Linear;
FH Key
FT source
PT source
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AUTHORS
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George, J.F. Jr. and Schroeder, H.W. Jr.
George, J.F. Jr. and Schroeder, H.W. Jr.
Developmental regulation of D beta reading frame and junctional diversity in T cell receptor-beta transcripts from human thymus
J. Immunol. 148 (4), 1230-1239 (1992)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      E09808.1 GI:22026437 JP 1995213280-A/26.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GenBank staff at the National Library of Medicine created this entry [NCBI gibbsq 81367] from the original journal article.

Location/Qualifiers
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Similarity 100.0%; Pred. No. 4.2e+04;
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                                                                                                                                                                                                /organism='Artificial sequences'.
                                                                                                                                     /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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/mol_type="mRNA"
/db_xref="taxon:9606"
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                                                       2.8%; Score 14; DB
100.0%; Pred. No. 4.2
tive 0; Mismatches
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CQ623100/c
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AR214531/c
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                                 Sequence 7840 1
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CQ623100.1 GI:
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Sequence 7839 from Patent
CQ623099
CQ623099.1 GI:41673317
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                                                                                                                                                                                                                                                                                                                                      Gu,Y., Ji,Y.,.Penn,S.G., Hanzel,D.K., Rank,D.R., Chen,W. and Shannon,M.E.
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                                                                      CQ623100
                                                                                                                                                                                                                                                                                                                Myosin-like gene expressed in human heart and Patent: WO 0192524-A 7839 06-DEC-2001;
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Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Patent: US 6407284-A 10 18-JUN-2002;
Location/Qualifiers
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Kudo,J., Takashima,Y. and Mine,S.
Method of resolving 2-oxobicyclo [3.1.0]
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Homo sapiens
Homo sapiens
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Similarity 100.0%;
13; Conservative (
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                                                                                                                                 CACCIGCIGCTIC 5
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                                                                                                                                                                                                                                      /organism="Homo sapiens"
/mol_type="unassigned DN
/db_xref="taxon:9606"
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                                    GI:41673318
            (human)
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JOURNAL
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Sequence 7842 from Patent
CQ623102
CQ623102.1 GI:41673320
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       Shannon, M.E.
Myosin-like gene expressed in human heart and Patent: WO 0192524-A 7842 06-DEC-2001;
                                              Gu,Y., Ji,Y., Penn,S.G., Hanzel,D.K., Rank,D.R., Chen,W. and
                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                  Homo sapiens (human)
Homo sapiens
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Sequence 7841 from Patent
CQ623101
CQ623101.1 GI:41673319
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Shannon, M.E.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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100.0%; Pred. No. 1.6e+05;
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   Patent: JP
TONEN CORP
OS Artifi
PN JP 2000
PD 04-APR
PF 28-SEP
PR
TOSHIY
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                                                                                              synthetic construct
synthetic construct
artificial sequences.

1 (bases 1 to 17)
Hayano, T., Hino, S. and Kajie, S.
Protein participating in oxidation reaction of organosulfur
compound and gene encoding the same
patent: JP 2000093180-A 17 04-APR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                 30 CACCTGCTGCTTC 42
                                                                                                                                                                                                                   E34099
E34099.1 GI:18624218
JP 2000093180-A/17.
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                                                                                                                                                                                                                                                                Protein participating in oxidation reaction compound and gene encoding the same.
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Sequence 7843 from Patent
CQ623103
CQ623103.1 GI:41673321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Myosin-like gene expressed in human heart and muscle Patent: WO 0102524-A 7843 06-DEC-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gu,Y., Ji,Y., Penn,S.G., Hanzel,D.K., Rank,D.R., Chen,W. Shannon,M.E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                       Artificial sequence JP 2000093180-A/17
                                                                                                                                                                                                                                                                                                                                                                                                                           2.6%; Score 13; DB 6; ilarity 100.0%; Pred. No. 1.6e+0 Conservative 0; Mismatches
TOSHIYA HAYANO, SANAE HINO, SHINICHI KAJIE
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/mol_type="unassigned DNJ
/db_xref="taxon:9606"
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/mol_type="unassigned DN/
/db_xref="taxon:9606"
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PC B09B3/00,(C12N15/00,C12R1:38)
PC CC Location/Qualifiers
FH Key 1..17
FT Bource /organism='Artificial sequence'.
                     Unclassified.

1 (bases 1 to 17)

Gu,Y., Ji,Y., Penn,S.G., Hanzel,D.K., Rank,D.R., Shannon,M.E.
Shannon,M.E.
Polynucleotide encoding a human myosin-like polyr predominantly in heart and muscle Patent: US 6686188-A 7840 03-FEB-2004;
                                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 17)
Gu,Y., Ji,Y., Penn,S.G., Hanzel,D.K., Rank,D.R., Chen,W.
Shannon,M.E.
Polynucleotide encoding a human myosin-like polypeptide e predominantly in heart and muscle
Patent: US 6686188-A 7839 03-FEB-2004;
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larity 100.0%; Pred. No. 1.6e+05;
Conservative 0; Mismatches 0;
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                                                                                                                                                                                                                                                                                                                                       /organism="unknown"
/mol_type="genomic DNA"
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15 CACCTGCTGCTTC 3
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Gu,Y., Ji,Y., Penn,S.G., Hanzel,D.K., Rank,D.R.,
Shannon,M.E.
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AR464164
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Gu,Y., Ji,Y., Penn,S.G., Hanzel,D.K.,
                                                                                                                                            Polynucleotide encoding a human myosin-like predominantly in heart and muscle Patent: US 6686188-A 7842 03-FEB-2004; Location/Qualifiers
                                                                                                                                                                                                                                                                                             AR464165
Sequence 7842 from patent
AR464165
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/mol_type="genomic
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/mol_type="genomic
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100.0%; Pr/
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A26384
A26384.1
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                      probe no.2.
                                   A26384
                                                                                                                                                                                                                                                                        Patent: WO 03040369-A 5772 15-MAY-2003; Molecular Engines Laboratories (FR)
                                                                                                                                                                                                                                                                                                             sequences involved in tumoral suppression, apoptosis and/or viral resistance phenomena
                                                                                                                                                                                                                                                                                                                           Telerman, A., Amson, R. and Tuijnder, M. Sequences involved in tumoral suppres
                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 5772 from Patent
AX762451
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 (bases 1 to 17)
Gu,Y., Ji,Y., Penn,S.G., Hanzel,D.K., Rank,D.R., Chen,W.
Shannon,M.E.
                                                                                                                                                                                                                                                                                                     medicines
                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Polynucleotide encoding a human myosin-like polypeptide expressed predominantly in heart and muscle Patent: US 6686189-A 7843 03-FEB-2004;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 7843 from patent US
AR464166
                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens (human)
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Unclassified.
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GI:904941
                                                                                                                                                                                                             /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="unknown"
/mol_type="genomic DNA"
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A89850
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A87883
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               1 (bases 1 to 18)

Brysch, W.D. and Schlingensiepen, K.D.
An antisense oligonucleotide preparation
Patent: EP 0856579-A 31 05-AUG-1998;
BIOGNOSTIK GES (DE)
Location/Qualifiers
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                                                                                                                       unidentified
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Brysch, W. and Schlingenslepen, K.
ANTISENSE OLIGONUCLECTIDE PREPARATION METHOD Patent: WO 9833904-A 31 06-AUG-1998;
BIOGNOSTIK GES (DE); BRYSCH WOLFGANG (DE)
Location/Qualifiers
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Patent: WO 9211289-A 10 09-JUL-1992;
Location/Qualifiers
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/organism="unidentified"
                                                                                                                                                               GI:6738364
                                                                                                                                                                                                                                                                                                                                                                                           /organism="unidentified"
/mol_type="unassigned DNA"
/db_xref="taxon:32644"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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AR300017/c
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AR297846/c
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Best Local Similarity 100.0%; Pred. No. 1.6e+05;
Matches 13; Conservative 0; Mismatches 0;
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Cohen, D., Chumakov, I. and Blumenfeld, M.
Biallelic markers for use in constructing
disequilibrium map of the human genome
Patent: US 6537751-A 9581 25-MAR-2003;
                                                                                                                                                                                            1 (bases 1 to 18)
Cohen, D., Chumakov, I. and Blumenfeld, M.
Biallelic markers for use in constructing
disequilibrium map of the human genome
Patent: US 6537751-A 11752 25-MAR-2003;
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Sequence 11752 from patent
AR300017
AR300017.1 GI:31687301
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                                                                             Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mol_type="unassigned DNA"
/db_xref="taxon:32644"
                                                                                                                                          /mol_type="genomic DNA"
                                                                                                                                                         organism="unknown"
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/mol_type="genomic DNA"
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100.0%; Pred. No. 1.6e+05;
ive 0; Mismatches 0;
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AX111590
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AX111589/c
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AX111597/c
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AX111597 18 bp
Sequence 27 from Patent WO0123561.
AX111597
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AX111589
AX111589.1 GI:13927870
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                                                                                                                                                                                                                                                                                     Patent: WO 0123561-A 20 05-APR-2001; Curagen Corporation (US)
                                                                                                                                                                                                                                                                                                                                                                                              synthetic construct synthetic construct
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AX111590
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Patent: WO 0123561-A 19 05-APR-2001;
Curagen Corporation (US)
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/mol_type="unassigned DNA"
/mol_type="unassigned DNA"
/db xrefe"ttaxon:32630"
/noEe="Chemically Synthesized"
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/mol_type="unassigned DNA"
/mol_tyse="caxon:32630"
/mote="Chemically Synthesized"
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100.0%; Pred. No. 1.6e+05;
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An antisense oligonucleotide
BD065396
BD065396.1 GI:22610999
JP 2001511000-A/31.
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13; Conserv
     unclassified
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Sequence 28 from Patent
AX111598
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/mol_type="unassigned DNA"
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/note="Chemically Synthesized"
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/mol_type="unassigned DNA"
/db xref="caxon:32630"
/noTe="Chemically Synthesized"
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100.0%; Pred. No. 1.6e+05;
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31-JAN-1997 EP 97101531
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                                                                                                                                                                                              /organism="Rattus norvegicus"
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/db_xref="taxon:10116"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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         GI:21066427
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Bilbe,G., Kinnunen,A. and Koenig,J.I.
Genes associated with schizophrenia, adhd and bipolar disorders
Patent: WO 2004007762-A 17 22-JAN-2004;
Novartis AG (CH); Novartis Pharma GmbH (AT); University of Mary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 (bases 1 to 18)
Schlingensiepen,K.H. and Brysch,W.
An antisense Oligonucleotide preparation method
Patent: JP 2001511000-A 31 07-AUG-2001;
BIOGNOSTIK GESELLSCHAFT FUR BIOMOLEKULARE DIAGNOSTIK MBH
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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Sequence 2 from Patent WO02068685.
AX670675
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          (bases 1 to 20)
Skolnick,M.H., Goldgar,D.E., Miki,Y.,
Harshman,K.D., Shattuck-Eidens,D.M., 7
and Futreal,P.Andrew.
                                                                     Unknown
                                                                                                                        AR004675
Sequence 4 from patent
AR004675
                                                                                                                                                                                                                                                                                                                                                                                                 Patent: WO 02068685-A 2 06-SEP-2002; Cytogenetic DNA Services Ltd (GB)
                                                                                                                                                                                                                                                                                                                                                                                                                                     Levett, L.J. and Liddle, S. Diagnostic test for the d
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/mol_type="unassigned DNA"
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100.0%; Pred. No. 1.6e+05;
                                                                                                                                                                                                                                                          2.6%; Score 13; DB 6; L6
L00.0%; Pred. No. 1.6e+05;
Lve 0; Mismatches 0;
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5747282.
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                       , Swenson, J., Kan
Tavtigian, S.V.,
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                                                                                                                                                linear
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                                    Kamb, A.
                                                                                                                                                PAT 04-DEC-1998
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                       Wiseman, R.W.
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AR008161/c
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l (bases 1 to 20)
l (bases 1 to 20)
Skolnick,M.H., Goldgar,D.E., Miki,Y., Swenson,J., Kamu,A.,
Harehman,K.D., Shattuck-Eidens,D.M., Tavtigian,S.V., Wiseman,R.W.
and Futreal,P.Andrew.
170-linked breast and ovarian cancer susceptibility gene
Patent: US 5753441-A 4 19-MAY-1998;
Location/Qualifiers
Location/Qualifiers
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Sequence 4 from patent US
AR008161
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17Q-linked breast and ovarian cancer susceptibility gene Patent: US 5747282-A 4 05-MAY-1998; Location/Qualifiers
                                                                                                                       1 (bases 1 to 20)
Skolnizk,M.H., Goldgar,D.E., Miki,Y.,
Harshman,K.D., Shattuck-Eidens,D.M., '
and Futreal,P.Andrew.
17q-liked breast and ovarian cancer |
Patent: US 6162897-A 4 19-DEC-2000;
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AR136944
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Similarity 100.0%; Pred. No.
13; Conservative 0; Mismatc
                                                                                                                                                                                                                                Unknown.
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                                                                                                           Location/Qualifiers
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/mol_type="unassigned
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                                                                                  organism="unknown"
                                                                    /mol_type="unassigned
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2.6%; but
100.0%; Pre
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100.0%; Pr
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Tavtigian, S.V.,
                                                                                                                                     susceptibility gene
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J., Wiseman,R.W.
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176945/c
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E12085/c
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PN JP 1996242897-A/1
PD 24-SEP-1996
PF 07-MAR-1995 JP 1995074616
PF 07-MAR-1995 JP 1995074616
PI IMAI HIDEKI PC C1201/68, COTH21/04, C12N15/09; CC strandedness: Single; CC typology: Linear; CC hypothetical: NO; FH Key Location/Qualifiers FT Source 1.20
FT source 1.20
FT source 1.20
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                                                                                                                                                              Linked breast and ovarian cancer susceptibility gene Patent: US 5693473-A 4 02-DEC-1997; Location/Qualifiers
                                                                                                                                                                                                                                                                             1 (bases 1 to 20)
Shattuck-Eidens, D.M., Simard, J., Durocher, F.,
                                                                                                                                                                                                                                                                                                                                             Unclassified.
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JP 1996242897-A/1.
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PCR primer for amplification
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                                                                                 /organism="unknown"
/mol_type="unassigned
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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RESULT 47
AR300298
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I80940/c
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 AR300298
Sequence
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                                                                                                                                                                                                                  Unclassified.

1 (bases 1 to 20)

1 (bases 1 to 20)

Skolnick,M.H., Goldgar,D.E., Miki,Y., Swenson,J., Kamb,A.,

Harshman,K.D., Shattuck-Eidens,D.M., Tavtigian,S.V., Wiseman,R.W.

and Futreal,P.Andrew.

17q-linked breast and ovarian cancer susceptibility gene

17q-linked breast and ovarian cancer susceptibility gene
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181036
181036.1
                                                                                                                     l Similarity
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Shattuck-Eidens, D.M., Simard, J., Durocher, F.,
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100 from patent US 6537775.
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                                                                                                                                                                          /organism="unknown"
/mol_type="unassigned
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/mol_type="unassigned DNA"
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                                                                                                               20 bp
Sequence 2 from Patent W003054230.
AX798039 AX798039.1 GI:37604329
                                           Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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13; Conserv
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Hampson, G.N.

Detection of disease due to abnormal oestrogen levels Patent: WO 03054230-A 2 03-JUL-2003;
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Patent: WO 02072882-A 5002 19-SEP-2002;
OGHAM GmbH (DE)
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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AX613977
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Location/Qualifiers
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AR300298.1
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/mol_type="unassigned DNJ
/db_xref="taxon:9606"
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/mol_type="genomic DNA"
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and therapeutic
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OS Homo sapiens (human)
PN JP 2001346593-A/3
PD 18-DEC-2001 JP 200111
                                        432
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ROGER W WISEMAN,
PI ANDREW P FUTE
PC C12N15/09,C12
PC 00,C12N5/00
CC Strandedness:
CC Topology: Lit.
CC Genes sensiti
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Genes sensitive to 17q-chained i
BD105585
BD105585.1 GI 22651159
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PR 12-AUG-1994 US 08/289221,02-SEP-1994 US 08/SEP-1994 US 08/SEP-1994 US 08/SEP-1994 US 08/SEP-1995 US 08/409305,07-JUN-1995 US 08/409305,07-JUN-1995 US 08/487002
PI MARK H SKOLNICK,DAVID E GOLDGAR,YOSHIO MIKJALEXANDER KAMB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Genes sensitive to 17q-chained breast cancer and ovarian cancer Patent: JP 2001346593-A 3 18-DEC-2001; MYRIAD GENETICS INC.UNIVERSITY OF UTAH RESEARCH FOUNDATION, THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 20)
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142
             CATTCCTGCTTCC 4
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C12N15/09,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12Q1/68,C12N15/
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                                                                                                                                                                                                                                                                                       Strandedness: Single;
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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/db_xref="taxon:9606"
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J. 1.6e+05;
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J. 1.6e+05;
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AX114821/c
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AX114695/c
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Sequence 106 from Patent
AX118750
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13; Conserv
            synthetic construct
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Agent for treating cephalic pain
Patent: WO 0128539-A 107 26-APR-2001;
GLAXO GROUP LIMITED (GB)
Location/Qualifiers
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Sequence 107
AX114695
                                        AX118750.1
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Therapy of cephalic pain
Patent: W0 0128540-A 107 26-APR-2001;
GLAXO GROUP LIMITED (GB)
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                                                                                                                                               TTCTGCCACCTGT 1
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ilarity 100.0%;
Conservative (
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mol_type="unassigned DNA"

db_xref="taxon:32630"
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/mol_type="unassigned DNA"
/db xref="taxon:32630"
/note="PCR Primer"
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                                       GI:14035701
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Pred. No. 1.6e+05;
0; Mismatches 0;
                                                                21 bp
WO0129255
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WO0128539.
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WO0128540.
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AX644766/c
LOCUS
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ACCESSION
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AX128167/c
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Patent: WO 0233121-A 104 25-APR-2002;
GLAXO GROUP LIMITED (GB)
Location/Qualifiers
                                              Hosford,D. and Purvis,I.J.
Test for the diagnosis of diabetes and compounds for
                                                                                           synthetic construct synthetic construct artificial sequences.
                                                                                                                                                                 Sequence 104 from Patent AX644766
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Cephalic pain susceptibility marker
Patent: WO 0129256-A 106 26-APR-2001;
GLAXO GROUP LIMITED (GB)
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Sequence 106 from Patent WO0129256.
AX128167
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Diagnostic test for cephalic pain
Patent: W 0129255-A 106 26-APR-2001;
GLAXO GROUP LIMITED (GB)
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Similarity 100.0%; Pred. No. 1.6e+05;
13; Conservative 0; Mismatches 0;
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/db_xref="taxon:32630"
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WO0233121.
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PAT 27-FEB-2003

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AR252249
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KEYWORDS
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AX815823
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  Query Match
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Best Local S
Matches 13
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TITLE
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Sequence 78
AX815823
                                                                                              1 (bases 1 to 22)
Eagles, P.A.M. and Zheng, R.Q.
Inhibition of cytokine production
Patent: US 6476214-A 2 05-NOV-2002;
Location/Qualifiers
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13; Conserv
                                                                                                                                                                                                                               Sequence 2 from patent US AR252249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hardge, T., Schellander, K. and Wimmers, K. Genetic markers for the diagnosis of the expression of nipples in pets, breeding animals and domestic cattle Patent: WO 03066891-A 78 14-AUG-2003; Foerderverein Biotechnologieforschung der deutschen Schweineproduktion e.V. (DE)
                                                                                                                                                                 Unknown.
Unclassified.
                                                                                                                                                                                            Unknown.
                                                                                                                                                                                                                   AR252249.1
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Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae; Sus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sus scrofa
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              Similarity
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  Conservative
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llarity 100.0%;
Conservative
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/mol_type="unassigned"
/db_xref="taxon:9823"
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|/organism="synthetic construct"
|/mol_type="unassigned DNA"
|/db_xref="taxon:32630"
|/note="PCR Primer"
                                                              /organism="unknown"
/mol_type="genomic
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from Patent WO03066891.
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              2.6%;
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Score 13; DB 6; Lo
Pred. No. 1.6e+05;
0; Mismatches 0;
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Pred. No. 1.6e+05;
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                       RESULT 60
AX224699
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AUTHORS
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KEYWORDS
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KEYWORDS
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AX003054
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Best Local Similarity
Matches 13; Conserv
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AX224699
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AX224699
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Sequence
AR489226
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 2 from Patent WO9937760.
AXO03054
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Inhibition of cytokine production
Patent: WO 9937760-A 2 9-JUL-1999;
EAGLES PETER ANTHONY MINTER (GB); ZI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent: US 6709860-A 5 23-MAR-2004;
Location/Qualifiers
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Enerback, S. and Carlsson, P.
Animal model
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             46
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/mal_type="unassigned DNA"
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/noEe="Human interleukin-4 promoter"
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tive 0; Mismatches
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WO0161021
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BD129696
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KEYWORDS
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                                                                                                                                         Query Match
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Best Local Similarity
Matches 13; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

E 1 (bases 1 to 22)

S Eagles, P.A.M. and Zheng,R.Q.
Inhibition of cytokine production
Patent: JP 2002500883-A 2 15-JAN-2002;

BTG INTERNATIONAL LTD
OS Homo sapiens (human)
PN JP 2002500883-A/2
PN J
                                         127 TGCCACCTGTACT 139
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A61P19/02,
PC A61P29/00
interleukin-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Inhibition of cytokine production. BD129696
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Cucumis melo
Cucumis melo
Eukaryota, Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
rosids; eurosids I; Cucurbitales; Cucurbitaceae; Cucumis.
                                                                                                                                               Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Transgenic plants having resistance to a fungal disease Patent: WO 0161021-A 46 23-AUG-2001;
Bar Ilan University (IL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cohen, Y. and Kenigbuch, D.
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TGCCACCTGTACT 19
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                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A61P29/00, A61P43/00//A61K38/00, C12N15/00, A61K37/02 CC
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                                                                                                                                                                                                                                                  /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                /organism='Homo
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/db_xref="taxon:3656"
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                                                                                                     2.6%; Score 13; DB
100.0%; Pred. No. 1.6
tive 0; Mismatches
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o. 1.6e+05;
0;
                                                                                                     1.6e+05;
thes 0;
                                                                                                                                                                 DB 6;
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                                                                                                                                                                                                                                                                                                                                                                                           sapiens (human)'.
                                                                                                                                                     Length 22;
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KEYWORDS
SOURCE
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AR036150/c
LOCUS
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ACCESSION
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DEFINITION
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BD234236/c
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Best Local Similarity
Matches 13; Conserv
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                                                                             JOURNAL
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                                              source
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                                                                                                                                                                                                                                                                                                                                                                                         405
                                                                                                                                                                                                                                                                                                                                                            13
                                                          Patent: US 5871992-A 8 16-FEB-1999;
Location/Qualifiers
                                                                                                     1 (bases 1 to 24)
Teebor,G.W. and Hilbert,T.P.
Mammalian endonuclease III, and diagnostic
                                                                                                                                                                                                                                           Sequence 8 from patent US AR036150
                                                                                                                                                                         Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A61P29/00,
PC CO7K14/47,CO7K16/18,C12N1/15,C12N1/19,C12P21/02,C12Q1/68,
PC G01N33/53,C12N15/00,C12N5/00,A61K37/02
PC Brimer Location/Qualifiers
PH Key Location/Qualifiers
FT Source 1.23
FT fource 1.23
                                                                                                thereof
                                                                                                                                                             Unclassified.
                                                                                                                                                                                              Unknown.
                                                                                                                                                                                                                           AR036150.1
                                                                                                                                                                                                                                                                        AR036150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATP-binding cassette genes and proteins for lipid disorders and inflammatory diseases Patent: JP 2002525111-A 46 13-AUG-2002; BAYER AKTIENGESELLSCHAFT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 23)
Schmitz,G. and Klucken,J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BD234236 23 bp DNA linear PAT 17-JUL-20 ATP-binding cassette genes and proteins for diagnosis and remedy lipid disorders and inflammatory diseases.

BD234236
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JP 200252511-A/46
13-AUG-2002
21-SEP-1999 JP 2000572359
25-SEP-1998 US 60/101706
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C12N15/09, A61K38/00, A61K45/00, A61K48/00, A61P3/06, A61P9/10,
          organism="unknown"
/mol_type="unassigned DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism='Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                           GI:5952818
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100.0%; Pred. No. 1.6e+05;
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                                                                                                        therapeutic uses
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Query Match Best Local Similarity

2.6%;

Score 13; Pred. No.

DB 6; L 1.6e+05;

Length

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BD183262
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FEATURES
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AR090647
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AR084729/c
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Matches 13
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Best Local
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BD183262 24 bp DNA 1 Method for estimating genotype of fertility against rice BT type male sterile cytoplasm. BD183262 BD183262.1 GI:31875462
                                                                                                            σ
                                                                                                                                                                                                                                                     l (bases 1 to 24)
Chenchik,A., Jokhadze,G. and Bibilashvilli,R.
Methods of assaying differential expression
Patent: US 5994076-A 767 30-NOV-1999;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 (bases 1 to 24)
Rio,M.-C., Tomasetto,C., Basset,P. and Byrne,J.
Isolated nucleic acid molecules useful as leukemia markers breast cancer prognosis and encoded polypeptides
Patent: US 5981218-A 53 09-NOV-1999;
                                                                                                                                                                                                                                                                                                                                Unknown
                                                                                                                                                                                                                                                                                                                                                                                Sequence 767
AR090647
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                                                                                                                                                                                                                                                                                                                  Unclassified.
                                                                                                                                                                                                                                                                                                                                            Unknown.
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                                                                                                                              AGCCACGCCCTGT 28
                                                                                                            AGCCACGCCCTGT 18
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llarity 100.0%;
Conservative
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/mol_type="unassigned
                                                                                                                                                                                                                  /organism="unknown"
/mol_type="unassigned
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                                                                                                                                                         Score 13; DB; Pred. No. 1.6
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                                   of fertility restoration
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J. 1.6e+05;
O;
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J. 1.6e+05;
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RESULT 68
AR259836
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ISM synthetic construct

artificial sequences.

CE 1 (bases 1 to 24)

RS Komori.T., Vanamoro,T., Nitta,N. and Takemori,N.

RS Komori.T., Vanamoro,T., Nitta,N. and Takemori.N.

Method for estimating genotype of fertility restoration gene site

against rice BT type male sterile cytoplasm

AL Patent: JP 2002345485-A, 1 03-DEC-2002;

JAPAN TOBACCO INC, SYNGENTA LTD

OS ARTIFICIAL Sequence

PN JP 2002345485-A/1

PD 03-DEC-2002

PF 17-AUG-2001 JP 2001247600

PF 17-AUG-2001 JP 2001247600

PF 17-AUG-2001 JP 2001247600

PF 17-AUG-2001 JP 2001247600

OC 01190Nucleotide primer for amplification

of R1877 EcoRI marker

CC sequence.

FH Key Location/Qualifiers

FT SOURCE /organism='Artificial Sequence'.
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Sequence
AR197682
           Sequence
AR259836
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                                                                                                                                               13;
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1. (bases 1 to 24)

1. (bases 1 to 24)

Chenchik, A., Jokhadze, G. and Bibilashvilli, R.

Methods of assaying differential expression

Patent: US 6352829-A 767 05-MAR-2002;

Location/Qualifiers
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/mol_type="unassigned
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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 GI:27310347
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                      from patent
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100.0%; Pred. No.
tive 0; Mismatcl
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                             Lawn,R.M., Wade,D., Oram,J.F. and Garvin,M. Atp binding cassette transporter protein at Patent: WO 0078971-A 27 28-DEC-2000; CV THERAPEUTICS, INC. (US)
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Regulation with binding cassette transporter protein abc1
Patent: WO 0078972-A 27 28 DEC-2000;
CV THERAPEUTICS, INC. (US)
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Sequence 27 from Patent
AX060739
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Chenchik,A., Jokhadze,G. and Bibilashvilli,R.
Methods of assaying differential expression
Patent: US 6489455-A 767 03-DEC-2002;
Location/Qualifiers
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organism="synthetic construct"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/noTe="ABC1 sequencing primer"
                      location/Qualifiers
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/mol_type="genomic DNA"
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BD141710
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Best Local
artificial sequences.

RE (bases 1 to 24)

Komori,T., Yamamoto,T., Nitta,N. and Takemori,N.

RE Komori,T., Yamamoto,T., Nitta,N. and Takemori,N.

RE Komori,T., Yamamoto,T., Nitta,N. and Takemori,N.

RE type male strimating genotype of fertility recovery locus for rice

BT type male sterile cytoplasm

AL Patent: WO 0214506-A 1 21-FEB-2002;

JAPAN TOBACCO INC,SYNCENTA LTD,TOSHIYUKI KOMORI,TOSHIO YAMAMOTO,

NAOTO NITTA,NAOKI TAKEMORI

OS Artificial Sequence
PN WO 0214506-A)

PD 21-FEB-2002

PF 16-AUG-2001 WO 2001JP007052

PF 17-AUG-2000 JP 00P 247204

PI TOSHIYUKI KOMORI,TOSHIO YAMAMOTO,NAOTO NITTA,NAOKI TAKEMORI PC

C12N15/09,C12Q1/68//A01H1/00

CC Oligonucleotide primer for amplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local
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Method of estimating
BT type male sterile
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/mol_type="unassigned DNA"
/db xrefs"ttaxon:32630"
/note="oligonucleotide primer"
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/db_xref="taxon:32630"
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artificial sequences.

E 1 (bases 1 to 25)

S Arata, Y., Nakajima, H. and Mukumoto, F.

Protein capable of catalyzing dehydrogenation reaction, ger

encoding said protein and use thereof

Patent: JP 2003024072-A 7 28-JAN-2003;

Sumitomo Chemical Co Ltd

OS Artificial Sequence

PN JP 2003024072-A/7
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1 (bases 1 to 25)

Sonigo,P., Brechot,C. and Courgnaud,V.

Sonigo,P., Brechot,C. and Courgnaud,V.

Oligonuclectide sequences for amplification of type HIV-2 and siv retroviruses genomes and their application to in-vitro diagnostic of infections caused by these viruses

Patent: BP 0404625-A 12 27-DEC-1990;

INSTITUT PASTEUR; INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE (INSERM)
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PF 06-JUL-2001 JP 2001205903
PI 'Syuto arata,hiroki nakajima,fujio mukumoto
CC Designed oligomucleotide primer for PCR
FH Key Location/Qualifiers.
                                                                                                                                                                                                                                              Gu,Y., Ji,Y., Penn,S.G., Hanzel,D.K., Rank,D.R., Chen,W. and Shamnon,M.E. M.B. Shamnon,M.E. Myosin-like gene expressed in human heart and muscle Patent: WO 0192524-A 12731 06-DEC-2001;
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Sequence
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Thompson, J., Mcswiggen, J., Mckenzie, T.,
                                                    Homo sapiens
Eukaryota; M
                                                                                                                                                                                                                                                                                                         l Similarity
15; Conser
                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                   Homo sapiens
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Patent: WO 0211674-A 904 14-FEB-2002;
RIBOZYME PHARMACEUTICALS, INC. (US) ;
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Method and reagent for the inhibition channel-1 (clca-1)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Location/Qualifiers
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904 from Patent WO0211674.
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/mol_type="unassigned RNA"
/db_xref="taxon:9606"
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/mol_type="unassigned RN/
/db_xref="taxon:9606"
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100.0%; Pr
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Patent: WO 0211674-A 905 14-FEB-2002; RIBOZYME PHARMACEUTICALS, INC. (US); Thompson, James (US)
  Telerman,A., Amson,R. and Tuijnder,M. Sequences involved in phenomena of tumour suppression, reversion, apoptosis and/or virus resistance and their medicines
                                                                Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                           Mus musculus (house mouse)
                                                                                                                                 AX727571.1 GI:30506914
                                                                                                                                             Sequence 5258 from Patent AX727571
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Patent: WO 0211674-A 903 14-FEB-2002;
RIBOZYME PHARMACEUTICALS, INC. (US) ; Sym
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Homo sapiens
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Similarity 100.0%; Pred. No. 1.5e+04;
14; Conservative 0; Mismatches 0;
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/mol_type="unassigned RNI
/db_xref="taxon:9606"
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                                                             Craniata; Vertebrata;
Sciurognathi; Muridae;
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                                                               Euteleostomi; Murinae; Mus.
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               Gold, D.P., Vainiene, M., Celnik, B., Wiley, S., Gibbs, C., Havandenbark, A.A. and Offner, H.
Characterization of the immune response to a secondary encephalitogenic epitope of basic protein in Lewis rats.
T cell. receptor V beta expression predominates in spinal infiltrating T cells
Infiltrating T cells
J. Immunol. 148 (6), 1712-1717 (1992)
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                                                                                                                                                           Rattus sp.
Eukaryota; Metazoa;
Mammalia; Eutheria;
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J. Immunol.
92176627
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Mammalia; Eutheria;
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:10090"
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Primates;
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thi; Muridae; Murinae;
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                                                AR064476.1
                                                                                        AR064476
                                                                                                                                                                                                                                                                                                                                                 Patent: WO 0211674-A 305 14-FEB-2002; RIBOZYME PHARMACEUTICALS, INC. (US); Thompson, James (US)
                                                                                                                                                                                                                                                                                                                                                                                         Method and reagent for the inhibition of calcium activated chloride channel-1 (clca-1)
                                                                                                                                                                                                                                                                                                                                                                                                                             Thompson, J., Mcswiggen, J., Mckenzie, T.,
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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llarity 100.0%;
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                                                                                                                                                                                                                                                                               /organism="Homo sapiens"
/mol_type="unassigned RN/
/db_xref="taxon:9606"
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/protein_id="AAB21736.1"
/db_xref="GI:246985"
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/codon_start=1
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                                                                Wirtz,R., Munnes,M. and Kallabis,H. Methods and compositions for the prediction, diprevention and treatment of malignant neoplasia Patent: EP 1365034-A 217 26-NOV-2003; Bayer HealthCare AG (DE).
                                                                                                                                               synthetic construct synthetic construct artificial sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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/organism="synthetic construct"
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/db_xref="taxon:32630"
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26 from Patent WO0206523.
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
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synthetic construct
artificial sequences.

CE 1 (bases 1 to 20)

CE 1 (bases 1 to 20)

Saito, A., Tamatsubo, K. and Adachi, K.

Genes and proteins participating in the upstream of degradation
passage of aromatic polycyclic compound
AL Patent: JP 2001245662-A 89 11-SEP-2001;
MARRINE BIOTECHNOLOGY INST CO LTD

OS Artificial Sequence
PN JP 2001245662-A/89

PD 11-SEP-2001

PF 03-MAR-2000 JP 200059523

PF 03-MAR-2000 JP 200059523

PF 03-MAR-2000 JP 200059523

PF 03-MAR-2000 JP 200059523

PG CINNISON CARJUAKI TAMATSUBO, KYOKO ADACHI
PC CINNISON CINNISON CO LTD

OC Description of Artificial Sequence: Synthetic primer KP234. FH

Key Increase A PR

Location/Qualifiers.
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                                             Patent: WO 2004056996-A 11 08-JUL-2004;
University of North Texas Health Science
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The escherichia coli carc gene and
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llarity 100.0%; F
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/db_xref="taxon:32630"
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    organism="synthetic construct"
/mol_type="unassigned DNA"
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tive 0; Mismatches
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Wilcox,E., Edwards,D.L., Schwab,G.E.,
Novel hybrid pesticidal toxins
Patent: EP 0349948-A1 17 08-NOV-1989;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                       Unknown
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Wilcox, E., Edwards, D.L., Schwab, G.E.,
Novel hybrid pesticidal toxins
Patent: EP 0340948-A1 6 08-NOV-1989;
Location/Qualifiers
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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CHIRON CORP

S Artificial Sequence
PN JP 2001510053-A/10
PD 31-JUL-2001
PD 31-JUL-2001
PR 18-JUL-1997 US 60/053066
PI XIN TAI CHEN,MEHDI GASMI,JIN KUANG YI,DOUGLAS J JOLLY PC C12N15/09,A61K48/00,C12N5/10,C12N7/00//A61K35/76,A61K38/00, PC A61K38/43,
PC C12N15/00,C12N5/00,A61K37/02,A61K37/465
PC Location/Qualifiers
PT SOURCE 1.28
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1 (bases 1 to 25)
Thompson, J.D. and Draper, K.G.
ErbB2/neu targeted ribozymes
Patent: US 5599704-A 67 04-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            JP 2001510053-A/10.
synthetic construct
synthetic construct
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1 (bases 1 to 28)
Chen, X. T., Gasmi, M., Yi, J.K.
Lentivirus vector
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BD076701.1 GI:22622304
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/mol_type="unassigned DNA"
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1 (bases 1 to 28)

Nishimura, M., Yaguchi, H., Naito, S. and Hiraoka, I.

Method for assaying an enzyme perticipating in conjugation and plucuronic acid in human beings, and probe and kit the Patent: UP 2002085066-A, 1 26-WAR-2002;

OTSUKA PHARMACEUTICAL FACTORY INC

OS Human UGT1 gene
PN JP 2002085066-A/1

PD 26-MAR-2002

PF 07-SEP-20^2

PF 07-SEP-20^2
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Jacobs, K., McCoy, J.M., LaVallie, E.R., Racie, L.A., Merberg, D.,
Treacy, M. and Spaulding, V.
Secreted proteins and polynucleotides encoding them
Patent: US 5723315-A 33 03-MAR-1998;
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 33 from patent US 
I89953
CAGCCCTGCATGC 13
                                                                             9.8%; Score 13; DB 6; L
llarity 100.0%; Pred. No. 5.5e+04;
Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Method for assaying an enzyme perticipating in conjugation with glucuronic acid in human beings, and probe and kit therefor FH Key
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MASUHIRO NISHIMURA,HIROSHI YAGUCHI,SHINSAKU NAITO,ISAO HIRAOKA
C12N15/09,C12Q1/25,C12Q1/68,G01N21/78,G01N33/50,G01N33/566, PC
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                                                                                                                                                                                 /organism="unknown"
/mol_type="unassigned
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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100.0%; Pred. No. 5.
tive 0; Mismatches
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S Merberg, D., Mccoy, J.M., Lavallie, E.R., Racie, L.A., Treacy, M., Spaulding, V. and Jacobs, K.

Secreted proteins and polynucleotides encoding them

Patent: JP 2001501460-A 24 06-FEB-2001;

GENETICS INSTITUTE INC
OS Unidentified
PN JP 2001501460-A/24
PD 06-FEB-2001
PP 22-AUG-1997 JP 1998511030
PR 23-AUG-1996 US 08/702344
PI DAVID MERBERG, JOHN M MCCOY, EDWARD R LAVALLIE, LISA A RACIE, MAURICE TREACY, TANGER CONTRACTS.
                                                                                                                       SM synthetic construct
artificial sequences.

1 (bases 1 to 29)

2 1 (bases 1 to 29)

3 Jacobs, K., Mccoy, J.M., Lavallie, E.R., Racie, L.A.C., Event artificial sand polynucleotides encoding them

Secreted proteins and polynucleotides encoding 
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PI CHERYL EVANS, DAVID MERBERG, MAURICE TREACY, VIKKI SPAULDING C12N15/09, C07K14/47, C12N5/10, C12P21/02, C12N15/00, C12N5/00 CC
                                                                                         PF 20-NOV-1998 JF 2000522129
PR 21-NOV-1997 US 08/976112,19-NOV-1998 US
KENNETH JACOBS,JOHN M MCCOY,EDWARD R LAVALLIE,LISA
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JP 2002504306-A/18.
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Secreted proteins and polynucleotides
BD138846
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PC C12N15/12,C12N5/10,C07K14/47,C12Q1/68,A61K38/17
Strandedness: Single;
CC Topology: Linear;
PH Key Location/Qualifiers
FT source 1. 29
PT forganism='Unidentified'.
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22-AUG-1997 JP 1998511030
23-AUG-1996 US 08/702344
23-AUG-1996 US 08/702344
DAVID MERBERG,JOHN M MCCOY,EDWARD R LAVALLIE,LISA A RACIE,
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llarity 100.0%; Pred. No. 5.5e+04;
Conservative 0; Mismatches 0;
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Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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CC: biotinylate
FH Key
FT misc_featur
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                                                 Gu,Y., Ji,Y., Penn,S.G., Hanzel,D.K., Rank,D.R., Chen,W. and Shannon,M.E.
Myosin-like gene expressed in human heart and muscle
Patent: WO 0192524-A 2182 06-DEC-2001;
                                                                                                                                                                                   CQ617442
Sequence 2182 from Patent
CQ617442
CQ617442.1 GI:41667660
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                                                                                                                  Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                              Homo sapiens
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                                                                                                                                                          Homo sapiens (human)
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/organism="Homo sapiens
                          Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/db_xref="taxon:9606"
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100.0%; Pr
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Primates;
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PAT 02-FEB-2004

Euteleostomi;; Homo.

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CQ617444/c
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CQ617443/c
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15 CCAGGACCTGCT 4
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                                                                                                                                                                                                                                                                                                        Sequence 2184 from Patent CQ617444 GI:41667662
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                                                                                                                                                           Myosin-like gene expressed in human heart and muscle Patent: WO 0192524-A 2184 06-DEC-2001;
                                                                                                                                                                                          Gu,Y., Ji,Y., Penn,S.G., Hanzel,D.K., Shannon,M.E.
                                                                                                                                                                                                                                                                 Homo sapiens (human)
                                                                                                                                                    Aeomica,
                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 h 9.1%; Score 12; DB 6; Length 17; Similarity 100.0%; Pred. No. 2e+05; 12; Conservative 0; Mismatches 0; Indels
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Similarity 100.0%; Pred. No. 2e+05;
12; Conservative 0; Mismatches 0; Indels
    9.1%; Scilarity 100.0%; F
                                                                        /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/db_xref="taxon:9606"
                  Score 12; DB 6; Pred. No. 2e+05;
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    0;
                               Length 17;
    Indels
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CQ617446/c
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Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                 Gu,Y., Ji,Y., Penn,S.G., Hanzel,D.K., Rank,D.R., Chen,W. and Shannon,M.E.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; E
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Beigelman,L., Burgin,A., Beaudry,A., Karpe
Matulic-Adamic,J., Sweedler,D. and Zinnen,
Synthetic ribonucleic acids with RNAse act
Patent: US 6528640-A 443 04-MAR-2003;
     1 (bases 1 to 17)
Gu,Y., Ji,Y., Penn,S.G.,
Shannon,M.E.
Polynucleotide encoding
                                                   Unknown.
Unclassified
                                                                                               AR458504
Sequence 2181 from patent
AR458504
AR458504.1 GI:42693561
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Beigelman, L., Burgin, A.B., Beaudry, A., Karpeisky, A.,
Matulic-Adamic, J., Sweedler, D. and Zinnen, S.
Oligoribonucleotides with enzymatic activity
Patent: US 6617438-A 442 09-SEP-2003;
Location/Qualifiers
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Sequence 442 from patent
AR398061
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                                                                                                                                                                                                                                                                                      /organism="unknown"
/mol_type="unassigned RNA"
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/mol_type="unassigned RNA"
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polypeptide
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Sequence
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Gu,Y., Ji,Y., Penn,S.G., Hanzel,D.K., Rank,D.R.,
Shannon,M.E.
                                                                                                                    Polynucleotide encoding a human myosin-like predominantly in heart and muscle Patent: US 6686188-A 2183 03-FEB-2004;
                                                                                                                                                      1 (bases 1 to 17)
Gu, Y., Ji, Y., Penn, S.G.,
Shannon, M.E.
                                                                                                                                                                                                                                                                                                                                                                      l Similarity
12; Conserv
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Patent: US 6686188-A 2181 03-FEB-2004;
Location/Qualifiers
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                                                                                                                                                                                        Unclassified.
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                     Conservative
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larity 100.0%; Pred. No. 2e
Conservative 0; Mismatches
                                                                                                                                                                                                                                                         2183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2182 from patent
                                                                         /organism="unknown"
/mol_type="genomic
                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="unknown"
/mol_type="genomic
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2e+05;
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2e+05;
                                                                                                                                                                Rank, D.R., Chen, W.
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                                        Length 17;
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RESULT 35
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AR458508/c
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AR458507/c
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    AR458509
Sequence
AR458509
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Sequence 2185 :
AR458508
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                                                                                                                                                                                                                                    Shannon, M.E.
Polynucleotide encoding a human myosin-like predominantly in heart and muscle predominantly in heart 2185 03-FEB-2004;
                                                                                                                                                                                                                                                                                 1 (bases 1 to 17)
GL,Y., Ji,Y., Penn,S.G., Hanzel,D.K., Rank,D.R., Chen,W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     l Similarity
12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Polynucleotide encoding a human myosin-like polypeptide expressed predominantly in heart and muscle Patent: US 6686189.A 2184 03-FEB-2004;
                                                                                                                                                                                                                                                                                                                        Unknown
                                                                                                                                                                                                                                                                                                                                       Unknown
                                                                                                                                                                                                                                                                                                                                                            AR458508.1
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Gu,Y., Ji,Y., Penn,S.G., Hanzel,D.K., Rank,D.R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AR458507
Sequence 2184 from patent
AR458507
                                                                                                                                                                                                                                                                                                              Unclassified.
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                                                                                                                                              Similarity
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                                                                                                     CCAGGACCTGCT 26
                                                                                     CCAGGACCTGCT 2
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llarity 100.0%; Pred. No. 2e+05;
Conservative 0; Mismatches
              2186
                                                                                                                                                                                         /organism="unknown"
/mol_type="genomic
                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="unknown"
/mol_type="genomic DNA"
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             patent
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Pred. No. 2e+
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              bp DNA 6686188.
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2e+05;
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                       PAT 20-FEB-2004
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AX579465/c
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AX579068/c
LOCUS
REFERENCE
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Best Local Similarity
Matches 12; Conserv
                                ORGANISM
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                                                                             Sequence
AX579465
      Homo sapiens
Eukaryota; Metazoa;
Mammalia; Eutheria;
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Sequence
AX579068
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                                           Homo sapiens (human)
                                                                  AX579465.1
                                                                                                    AX579465
                                                                                                                                                                                                                                                                                                                                     Thompson,J., Mcswiggen,J., Mckenzie,T., and Grupe,A.
Method and reagent for the inhibition of channel-1 (clca-1)
Patent: WO 0211674-A 906 14-FEB-2002;
RIBOZYME PHARMACEUTICALS, INC. (US); Symbosom
                                                                                                                                                                                                                                                                                                                         RIBOZYME PHARMACEUTICALS, Thompson, James (US)
                                                                                                                                                                                                                                                                                                                                                                                                                    Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi; Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Polynuclectide encoding a human myosin-like predominantly in heart and muscle Patent: US 6686188-A 2186 03-FEB-2004;
Location/Qualifiers
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Gu,Y., Ji,Y., Penn,S.G., Hanzel,D.K., Rank,D.R.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                        1303 from Patent
                                                                                                                                                                                                                                                                   /organism="Homo sapiens"
/mol_type="unassigned RN/
/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
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/mol_type="genomic
                                                                 GI:27648667
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100.0%;
       Chordata;
Primates;
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WO0211674
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       Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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2e+05;
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2e+05;
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PAT 10-JAN-2003

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PAT 10-JAN-2003

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REFERENCE
AUTHORS
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BD012743/c
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AUTHORS
TITLE
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DEFINITION
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AX731734/c
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Best Local (
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                                                                                                                                        A novel :
BD012743
synthetic construct
synthetic construct
artificial sequences.

1 (bases 1 to 18)
Seno, C. and Numata, M.
A novel frizzled family gene, 584
Patent: WO 0112808-A 16 22-FFB-2001;
CHUGAI RESEARCH INSTITUTE FOR MOLECULAR MEDICINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12;
                                                                                                             BD012743.1 GI:22092932
WO 0112808-A/16.
                                                                                                                                                                    BD012743
                                                                                                                                                                                                                                                                                                                                                                                                                  Patent: WO 03025175-A 3368 27-MAR-2003;
Molecular Engines Laboratories (FR)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequences involved in phenomena of tumour suppression, tumoreversion, apoptosis and/or virus resistance and their use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Telerman, A., Amson, R. and Tuijnder, M. Sequences involved in phenomena of tumour
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; E
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 3368 from Patent
AX731734
AX731734.1 GI:30511077
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Method and reagent for the inhibition of channel-1 (clca-1)
Patent: WO 0211674-A 1303 14-FEB-2002; RIBOZYME PHARMACEUTICALS, INC. (US) ; SyrThompson, James (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                             medicines
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens (human)
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ilarity 100.0%;
Conservative
                                                                                                                                                     frizzled family
                                                                                                                                                                                                                                                                                                                                                            /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Homo sapiens"
/mol_type="unassigned RNA"
/db_xref="taxon:9606"
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WO03025175
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  INC, CHIAKI SENO,
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RESULT 41
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KEYWORDS
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AUTHORS
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A51892
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unidentified unidentified unclassified.
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pN WO OIL
pD 18-AUG
PF 18-AUG
PR 18-AUG
PR CHIAKI
PC C12N15
C07K14/705,
PC C07K16
CC Descri
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                                                                Sequence 110 from Patent
A67354
A67354.1 GI:4756298
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Blakey, D.C., Davies, D.H., Dowell, R.I.,
Slater, Anthony, M. and Hennequin, L.F.
CHEMICAL COMPOUNDS CHEMICAL COMPOUNDS
Patent: WO 9620011-A 56 04-JUL-1996,
ZENECA LTD (GB)
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A51892
A51892.1 GI:2304640
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12; Conserv
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Key
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WO 0112808-A/16
22-FEB-2001
18-AUG-2000 WO 2000JP005552
18-AUG-2000 WO 2000JP005552
18-AUG-1999 JP 99P 232018
CHIAKI SENO, MARIKO NUMATA
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56 from Patent WO9620011.
                                                                                                                                                                                                                                                                                                                                                            /organism="unidentified"
/mol_type="unassigned DNA"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Pred. No.
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2e+05;
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PAT 07-OCT-1997

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DEFINITION
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AR085857
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KEYWORDS
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AUTHORS
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Best Local Similarity
Matches 12; Conserv
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                                                        1 (bases 1 to 19)
Taylorson, C.John., Eggelte, H.Johannes., Tarragona-Fiol, A., Rabin, B.Robert., Boyle, F.Thomas., Hennam, J.Frederick., Blakey, D.Charles., Marsham, P.Robert., Heaton, D.William., Davies, D.Huw., Slater, A.Michael. and Hennequin, L.Francois.
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                                                                                                                                                                                            Sequence 52
AR085857
                                         Patent: US 5985281-A 52 16-NOV-1999;
                                                                                                                              Unclassified.
                                                                                                                                            Unknown
                                                                                                                                                     Unknown.
                                                                                                                                                                               AR085857.1
                                                                                                                                                                                                                  AR085857
                                                                                                                                                                                                                                                                                                                                                                                                                                     Patent: WO 9835988-A 7 20-AUG-1998;
ZENECA LTD (GB); EDGE MICHAEL DEREK
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 19)
Edge, M.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            unidentified unidentified
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Sequence 7 from Patent WO9835988.
A87526
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COPLEY, C.G., Edge, M.D. and Emery, S.C.
MONOCLONAL ANTIBODY TO CEA, CONUUGATES COMPRISING SAID ANTIBODY,
AND THEIR THERAPEUTIC USE IN AN ADEPT SYSTEM
Patent: WO 9742329-A 110 13-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A87526.1 GI:6736175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                unclassified
                                                                                                                                                                                                                                                                                                                                      h 9.1%; Score 12; DB 6; Similarity 100.0%; Pred. No. 2e+05;
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                                                                                                                                                                                                                                                                                                                             Conservative
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                                                       compounds
     /organism="unknown"
                             Location/Qualifiers
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/mol_type="unassigned DNA"
/db_xref="taxon:32644"
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/mol_type="unassigned DNA"
/db_xref="taxon:32644"
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100.0%; Pred. No. 2e-
tive 0; Mismatches
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3 5985281.
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                                                               and Hennequin, L. Francois. Andre
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RESULT 46
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AR166750/c
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AR166750
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                                                                                                                                                                               Method for creating osmotic-pressure-tolerant plant
Patent: US 6281412-A 6 28-AUG-2001;
Location/Qualifiers
1. 20
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Sequence
I46738
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Murata, N.
                                                                                                                                                                                                                                                                      Unknown
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                                                                                                                                                                                                                                                                                                                                               AR166750
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      unclassified.

Unclassified.

(bases 1 to 19)

Mitsuhashi, M. and Cooper,A.

Method for detecting polynucleotides with polynucleotide probes identified based on Patent: US 5639612A, 717, 17-UN-1997;

Location/Qualifiers
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12; Conserv
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                                                                                                                                                           /mol_type="unassigned DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="unknown"
/mol_type="unassigned
                                                                                                                                                                       organism="unknown"
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100.0%;
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. 2e+05;
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2e+05;
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linear

PAT 17-OCT-2001

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Gaps

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Indels

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AR228892
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12; Conserv
w
                                                                                                                                      expression Patent: US
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Monia, B.P., Gaarde, W.A., Nero, P. and McKay, R.
Antisense modulation of p38 mitogen activated
                                                                                                                                                                                                                                            Sequence 99 from patent AR228892
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                                                                                                                                                                                        Unclassified.
                                                                                                                                                                                                    Unknown
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                    ACGITCIGCGCG 72
 ACGITCIGCGCG 14
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                                         Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C12N15/09, A61K31/711, A61P19/02, A61P29/00, A61P29/00, A61P37/06,
                                                                                                                          US 6448079-A 99 10-SEP-2002;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism='Artificial
                                                                                          /organism="unknown"
/mol_type="genomic DNA"
                                       9.1%; Score 12; DB
100.0%; Pred. No. 2et
tive 0; Mismatches
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RESULT 48
AR300685/c
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AR314724/c
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AR314616
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                                                                                                                                                                                                          12;
Unknown.
Unclassified.
1 (bases 1 to 20)
Griffais,R., Hoiseth,S.K., Zagursky,R.J., Metcalf,B.J., Peek,J.A.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2h 9.1%; Score 12; DB 6;
l Similarity 100.0%; Pred. No. 2e+05;
12; Conservative 0; Mismatches
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                                                                       Sequence 5261 from patent AR314724 AR314724.1 GI:31708150
                                                                                                                                                                                                                                                                                                         Chlamydia pneumoniae polynucleotides and Patent: US 6559294-A 5153 06-MAY-2003;
                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 20)
Griffais,R., Hoiseth,S.K., Zagursky,R.J.,
Sankaran,B. and Fletcher,L.D.
                                                                                                                                                                                                                                                                                                                                                                            Unknown.
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Sequence 5153 from patent
AR314616
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Unclassified.
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AR300685
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Freier, S.M.
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/mol_type="genomic |
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100.0%; Pred. No. 2e
tive 0; Mismatches
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CQ802500
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CQ802500
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AX511476
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                                                                                            Klippel-Giese, A., Kaufmann, J. and Schwarzer, R. Factor involved in metastasis and uses thereof Patent: WO 2004035615-A 13 29-APR-2004;
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Chlamydia pneumoniae polymucleotides and uses thereof
Patent: US 6559394A 5261 06-MAY-2003;
Location/Qualifiers
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larity 100.0%; Pred. No. 2e
Conservative 0; Mismatches
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  /note="antisense oligonucleotide"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="unassigned DNA"
/db xref="taxon.32630"
/noTe="Synthetic Oligonucleotide"
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/mol_type="genomic DNA"
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AR487575
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Sequence 59 from Patent W00077204.
AX057523
AX057523.1 GI:12310257
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                                                                                                                             Variant tlr4 nucleic acid and uses thereof Patent: WO 0077204-A 59 21-DEC-2000; University of Iowa Research Foundation (US)
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Han,H.-Q. and Kwak,K.
Human B3.alpha. ubiquitin ligase family
Patent: US 6706505-A 29 16-MAR-2004;
Location/Qualifiers
                                                                                                                                                                                                                                         Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unknown.
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                                                                                                                                                                              Forenz, E., Schwartz, D.A. and Schutte, B.C.
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                                                                              /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="genomic
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                                  Score 12;
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Pred. No. 2e+05;
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Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                    Length 21;
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                                                                                                                  Lorenz,
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RESULT 57
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AX740333/c
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Sequence 67 from Patent EP1300419.
AX740333
AX740333.1 GI:30523506
                                                                                                                                                                                                                                                    Antibody of human origin for inhibiting thrombocyte aggregation Patent: EP 1300419-A 67 09-APR-2003;
Affimed Therapeutics AG (DE)
Location/Qualifiers
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The Secretary of the Department of Health Location/Qualifiers
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AX179541
 CQ802965
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milarity 100.0%;
Conservative 0
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/mol type="genomic DNA"
/db xref="taxon:32630"
/noTe="Oligonucleotide"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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AX553601
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Sequence 5 from Patent WO02074946.
AX553601
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CQ802965
CQ802965.1 GI:47109960
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Patent: WO 02074946-A 5 26-SEP-2002;
NOVO NORDISK A/S (DK)
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synthetic construct
            artificial sequences
                                                               AX955623.1
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="unassigned DNA"
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/db_xref="taxon:32630"
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100.0%; Pred. No. 2e+05;
tive 0; Mismatches
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12; Conserv
                                                                                                       METHODS FOR PRODUCING MEMBERS OF SPECIFIC BINDING Patent: WO 9201047-A 82 23-JAN-1992;
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Synthetic PCR primer
A32959
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Oligonucleotide primer
A27365
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                                                organism="synthetic construct"
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                                                                                                Location/Qualifiers
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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Other publication AU 4832493 940426
Other publication JP 8501699T 960227
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Hawkins, R. E., Russell, S. J., Stevenson, F. K. and Winter, G. P. IMPROVEMENTS IN OR RELATING TO IMMUNE RESPONSE MODIFICATION PALENT: WO 9408008-A 14 14-APR-1994;
MEDICAL RES COUNCIL (GB)
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Sequence 14 from Patent W09408008.
A38170
A38170.1 GI:2294776
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Jespers, L.S.A., Winter, G.P., Baier, M. and Hoogenboom, H.R.J.

Production of chimeric antibodies - a combinatorial approach
Patent: WO 9306213-A 76 01-APR-1993;

CAMBRIDGE ANTIBODY TECH (GB); MEDICAL RES COUNCIL (GB)
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/mol_type="unassigned DNA"
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Blakey, D.C., Davies, D.H., Dowell, R.I.,
Slater, Anthony, M. and Hennequin, L.F.
CHEMICAL COMPOUNDS CHEMICAL COMPOUNDS
Patent: WO 9620011-A 43 04-UUL-1996;
ZENECA LTD (GB)
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Sequence 6 from Patent WO9620011.
A51842
           A51891
Sequence 55 from Patent
A51891
A51891.1 GI:2304639
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On Mar 27, 1999 this sequence version 1
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/mol_type="unassigned DNA"
/db_xref="taxon:32644"
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Blakey, D.C., Davies, D.H., Dowell, R.I., He
Slater, Anthony, M. and Hennequin, L.F.
CHEMICAL COMPOUNDS CHEMICAL COMPOUNDS
Patent: WO 9620011-A 55 04-JUL-1996;
ZENECA LTD (CB)
Other publication AU 4269796 960719.
Location/Qualifiers
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Sequence 6 from Patent WO9835988
A87525
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ZENECA LTD (GB); EDGE MICHAEL DEREK
Location/Qualifiers
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hes 12; Conservative 0;
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Taylorson, C. John., Eggelte, H. Johannes., Tarragona-Fiol, A., Rabin, B. Robert., Boyle, F. Thomas., Hennam, J. Frederick., Blakey, D. Charles., Marsham, P. Robert., Heaton, D. William., Davies, D. Huw., Slater, A. Michael. and Hennequin, L. Francois. Chemical compounds

Patent: US 5985281.A 39 16-NOV-1999;
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      l Similarity
12; Conserv
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Griffiths, A., David., Williams, S.Cameron., Waterhouse, P.Michael., Nissim, A., Winter, G.Paul., Johnson, K.Stuart. and Smith, A. John. Hammond.
Methods for producing recombinant vectors
Patent: US 5962255-A 554 05-OCT-1999;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12;
                                                                                                                                                                                                                                                          Sequence 39
AR085844
                                                                                                                                                                                               Unclassified.
                                                                                                                                                                                                          Unknown
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ilarity 100.0%;
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                                                            /organism="unknown"
/mol_type="unassigned
                                                                                                Location/Qualifiers
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
/db_xref="taxon:32644"
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AR085856/c
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pN JP 2003543830-A/22
pD 24-DEC-2002
pP 18-MAY-2000 JP 2000618429
pF 18-MAY-2000 JP 2000618429
pF 18-MAY-1999 EP 99201558.6
pI HENDRICUS RENERUS JACOBUS MATTHEUS HOOGENBOOM PC C12N15/09, COTAIN6/34, G01.N33/53, C12N15/00
CC Description of Artificial Sequence: Primer Location/Qualifiers
FT source 1...23
FT source 1...23
FT source 1...23
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Sequence
AR085856
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1 (bases 1 to 23)

Hoogenboom,H.R.J.M.

Novel Fab fragment libraries and method of using the same Patent: JP 2002543830-A 22 24-DEC-2002;
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Novel Fab fragment libraries
BD271430
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Taylorson, C. John., Eggelte, H. Johannes., Tarragona-Fiol, A., Rabin, B. Robert., Boyle, F. Thomas., Hennam, J. Frederick., Blakey, D. Charles., Marsham, P. Robert., Heaton, D. William., Davies, D. Huw., Slater, A. Michael. and Hennequin, L. Francois. Andre. Chemical compounds
Chemical compounds
Patent: US 595281-A 51 16-NOV-1999;
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                                                                       /organism="synthetic construct"
/mol_type="genomic DNA"
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                                                                                                                                /organism='Artificial
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Pred. No. 2e+05;
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E09118/c
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            I (bases 1 to 23)

E 1 (bases 1 to 23)

Kattou, T., Yamamoto, K., Nishioka, K. and Mizushima, Y.,

METHOD FOR DETECTING AND CLONING ANTISODY GENE
METHOD FOR DESILE978-A 13 09-MAY-1995;

L T T KENKYUSHO:KK

OS None

OC Artificial sequences.
PN JP 1995115978-A/13

PD 09-MAY-1995

PF 22-OCT-1993 JP 1993287628

PF 22-OCT-1993 JP 1993287628

PF 22-OCT-1993 JP 1993287628

PF KATOU TOMOHIRO, YAMAMOTO KAZUHIKO, NISHIOKA KUZUKI, PI

MIZUSHIMA YUTAKA,

PC C12N15/09,C12Q1/68,G01N33/53;

CC strandedness: Single;

CC topology: Linear;
FH Key Location/Qualifiers
FH Key Location/Qualifiers
FH Key /organism='Artificial sequences' FT

FT source 1. .23

PT source /organism='Artificial sequences' FT
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Synthetic oligonucleotides for
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Sequence 103 from Patent EP1433846.
CQ846582
                                                                                                                                                                                                                                                                                                                                                                          unidentified
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JP 1995115978-A/13.
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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100.0%; Pred. No. 2e+05;
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Job time : 564.527
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Hoogenboom,H.R.J.M., Baier,M., Jespers,L.
Production of chimeric antibodies - a com
Patent: US 5565322-A 76 15-OCT-1996;
Location/Qualifiers
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76 from patent US 5565332.
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AR197688 Sequence
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AR196704
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Zwick, M.G., Edington, B.E., McSwiggen, J.A., M. Guo, L., Skokut, T.A., Young, S.A., Folkerts, O. Nucleic acid encoding delta-9 desaturase Patent: US 6350934-A 1169 26-FEB-2002;

Location/Qualifiers
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1 (bases 1 to 18)

2 wick, M.G., Edington, B.E., McSwiggen, J.A., McGuo, L., Skokut, T.A., Young, S.A., Folkerts, O. Nucleic acid encoding delta-9 desaturase Patent: US 6350934-A 1167 26-FEB-2002;

Location/Qualifiers
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Miraglia,L., Bennett,C.Frank., Dean,N.
Antisense compositions and methods for
interleukin-1 receptor expression
Patent: US 5856099-A 1 05-JAN-1999,
Location/Qualifiers
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l Similarity 100.0%; Pred. No.
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Sequence 4808 from Patent
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Sequence 27 from patent
AR084538
AR084538.1 GI:10011309
                                                                                                                                                                                                                             Telerman,A., Amson,R. and Tuijnder,M.
Sequences involved in phenomena of tumour suppression, reversion, apoptosis and/or resistance to viruses and thereof as medicaments
Patent: WO 03025177-A 4808 27-MAR-2003;
Molecular Engines Laboratories (FR)
Location/Qualifiers
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Matson.R.S., Coassin,P.J., Rampal,J.B.
Oligonucleotide repeat arrays
Patent: US 5981185-A 27 09-NOV-1999;
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  AX452047
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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/mol_type="unassigned DNA"
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/mol_type="unassigned DN/
/db_xref="taxon:9606"
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100.0%; Pred. No. 1.4e+04;
Live 0; Mismatches 0;
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AX452047
AX452047.1 GI:21712050
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Miraglia, L., Bennett, C. Frank., Dean,
Antisense compositions and methods f
interleukin-1 receptor expression
Patent: US 5856099-A 2 05-JAN-1999;
1 (bases 1 to 20)
Griffais,R., Hoiseth,S.K., Zagursky,R.J.,
Sankaran,B. and Fletcher,L.D.
Chlamydia pneumoniae polynucleotides and v
                                                                Unknown
                                                                                                               Sequence 2554 from patent AR312017
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AR026495
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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Sequence 159 from Patent WO02059315.
AX686350
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AX354407
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Location/Qualifiers
                                                                                    Kennedy,G.C., Kang,S., Reinhard,C. and Jefferson,A.B. Polynucleotides related to colon cancer Patent: WO 0196523-A 53 20-DEC-2001; CHIRON CORPORATION (US)
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/organism="synthetic construct"
/mol_type="unassigned DNA"
/mb.xref="taxon:32630"
/noce="primer"
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/db_xref="taxon:32630"
/note="PCR Primer sequence"
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Direct Submission

Submitted (21-NOV-2002) Balzergue S., UMRGV, INRA/CNRS, 2 rue

Submitted (21-NOV-2002) Balzergue S., UMRGV, INRA/CNRS, 2 rue

Gaston Cremieux, 91057 Evry cedex, FRANCE

PCR was performed on DNA from transformants of Arabidopsis thaliana

plants from INRA (Versailles). The DNA fragment(s) resulting from

the PCR were directly sequenced from the left or the right border

to determine the genomic sequence flanking the insertion. T-DNA

derived sequences were removed. Information to order the

corresponding mutant line and a link to a database providing a

graphical display of the insertion site are available at

http://dbsgap.versailles.inra.fr/publiclines/. This sequence has

been generated in the framework of the French plant genomics

program 'Genoplante' (http://www.genoplante.com and

http://genoplante-info.infobiogen.fr).
                                              207 ATGAAAAATCAGGT 220
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Balzergue, S.
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EMBO Rep. 3 (12), 1152-1157 (2002)
22363535
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left border; T-DNA flanking sequence.

Arabidopsis thaliana (thale cress)

Arabidopsis thaliana
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Arabidopsis thaliana T-DNA flanking sequence, left border, clone
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                                                                                                                                                                                                                                                                                                                        /organism="Arabidopsis thaliana"
/mol_type="genomic DNA"
/cultivar="Waasillewskija"
/db_xref="taxon:3702"
                                                                                                                                                                                                           left border"
                                                                                                                                                                                                                              note="T-DNA flanking sequence
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L. .27
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100.0%; Pred. No. 1.9
ive 0; Mismatches
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RESULT 12 AX791502

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Sequence 190 from patent
AR033424
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13; Conserv
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Method for analysing a patient's predisposition to insulin-dependent diabetes, device and set of prime Patent: WO 0240711-A 11 23-MAY-2002;
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Helicobacter pylori
Bacteria, Proteobacteria, Epsilon
Helicobacteraceae, Helicobacter.
           Unknown.
                                                   AR033424.1
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 11 from Patent W00240711.
AX431364 1 GI:21656219
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Sequence 3966 from Pater
AX791502
AX791502.1 GI:32956949
   Unclassified.
                           Unknown.
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                                                                                                                                                                                                                                                              /organism="Homo sapiens"
/mol_type="unassigned DN
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:210"
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Pred. No. 5.6e+04;
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Draper, K.G.
Method and reagent for inhibiting hepatitis
Patent: US 5869253-A 190 09-FEB-1999;
                                                                                                                                                                                       unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 190
AR113246
                                                                                                                                                                                                               JP 2002512791-A/747.
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PAVCO,
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/mol_type="unassigned
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/mol_type="unassigned DNA"
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100.0%; Pr
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Blatt,I., Mcswiggen,J.A., Roberts,E., Pavco,P.A. and Macejak,D.
Enzymatic nucleic acid treatment of diseases or conditions related
to hepatitis C virus infection
Patent: JP 2002512791-A 747 08-MAY-2002;
RIBOZYME PHARMACEUTICALS INC
OS Hepatitis virus (hepatitis C virus)
PN JP 2002512791-A/747
PD 08-MAY-2002
PF 26-APR-1999 JP 2000545991
                                                                                                                                                                                                                                                                                                                                                                                      Enzymatic nucleic acid treatment of diseases or conditions related to hepatitis C virus infection.
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Method and reagent for inhibiting hepatitis
Patent: US 6132966-A 190 17-OCT-2000;
Location/Qualifiers
PF 26-APR-1999 JP 2000545991
27-APR-1998 US 60/083217,18-SEP-1998 US 60/100842
25-FEB-1999 US 09/257608,23-MAR-1999 US 09/274553 PI
LAWRENCE BLATT, JAMES A MCSWIGGEN, ELISABETH ROBERTS, PAMELA A
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BD104567/c
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I57653/c
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Nishida,M.
Nishida,M.
Nishida,M.
Kit and method for determining HLA type
Patent: WO 0192572-A 671 06-DEC-2001;
PATENT: WO 01925712-A 671 06-DEC-2001;
NISSHINBO INDUSTRIES INC, SYSTEM RESEARCH INC, HIDETOSHI INOKO, TAEKO
KAGIYA, TATSUO ICHIHARA, YOSHIYUKI MATSUMURA, SHOGO MORIYA, MICHIO
                                                                                            synthetic construct
synthetic construct
artificial sequences.
1 (bases 1 to 16)
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157653
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                                                                                                                                            BD104567.1 GI:22650141
WO 0192572-A/671.
                                                                                                                                                                    Kit and method for determining BD104567
                                                                                Inoko, H., Kagiya, T., Ichihara, T., Matsumura, Y.,
                                                                                                                                                                                           BD104567
                                                                                                                                                                                                                                                                                                                                                                                                       Enzymatic RNA molecule targeted against Hepatitis Patent: US 5610054-A 190 11-MAR-1997;
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Draper, K.G.
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A61K37/66,
C12N15/00
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/mol_type="genomic RNA"
/db_xref="taxon:32644"
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/mol_type="unassigned DNA"
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                          Homo sapiens (human)
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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Patent: WO 0188124-A 1960 22-NOV-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); GLAXO
                                                                                                        AX423625.1
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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WO 0192572-A/671
06-DEC-2001
01-JUN-2001 WO 2001JP004662
01-JUN-2000 JP 00P 164798
HIDETOSHI INOKO,TAEKO KAGIYA,TATSUO ICHIHARA,YOSHIYUKI
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C12Q1/68, C12M1/00, C12N15/09, G01N33/53
Description of Artificial Sequence: capture
Key Location/Qualifiers
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/mol_type="unassigned RNA"
/db_xref="taxon:9606"
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Sequence 210 from Patent
AX498903
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Sequence 1962 from Patent
AX423626
AX423626.1 GI:21527008
                            Human testis expressed patched like protein Patent: EP 1229046-A 210 07-AUG-2002;
                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                               Method and reagent for the inhibition of erg Patent: WO 0188124-A 1962 22-NOV-2001; RIBOZYME PHARMACEUTICALS, INC. (US); GLAXO (Location/Qualifiers
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Patent: WO 0188124-A 1961 22-NOV-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); GLAXO GROUP
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                                                         Zhan,J.
                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 212 from Patent
AX498905
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Sequence 211 from Patent
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Mammalia; Eutheria;
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h 4.1%; So Similarity 100.0%; F 13; Conservative 0;
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Similarity 100.0%; Pred. No. 5.7e+04;
13; Conservative 0; Mismatches 0;
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larity 100.0%; Pred. No. 5.7e+04;
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/db_xref="taxon:9606"
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Local Similarity 100.0%; Pred. No. 5.7e+04;
hes 13; Conservative 0; Mismatches 0;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                    Homo sapiens (human)
                                                                                         Sequence 103 from Patent AX530594
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                                                                                                                                                                                  4 GCGCGGAGGAGGG 16
                                                                          AX530594.1 GI:25252561
                                                                                                                 AX530594
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Patent: EP 1239051-A 102 11-SEP-2002;
Aeomica, Inc. (US)
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Sequence 102 from Patent EP1239051.
AX530593
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Mammalia; Eutheria; Primates;
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Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Patent: EP 1239051-A 101 11-SEP-2002;
Aeomica, Inc. (US)
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Similarity 100.0%; Pred. No. 5.
13; Conmervative 0; Mismatches
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Mammalia; Eutheria; F
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                                                            Human posh-like protein 1
Patent: EP 1239051-A 105 11-SEP-2002;
Aeomica, Inc. (US)
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Sequence 105 from Patent
AX530596
AX530596.1 GI:25252563
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Patent: EP 1239051-A 104 11-SEP-2002;
Aeomica, Inc. (US)
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Mammalia; Eutheria; Primates;
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Sequence 104 from Patent
AX530595
AX530595.1 GI:25252562
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Patent: EP 1239051-A 103 11-SEP-2002;
Aeomica, Inc. (US)
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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100.0%; Pred. No. 5.7e+04;
tive 0; Mismatches 0;
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A68426/c
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                                                                                                                                          Novel fissr-pcr primers and method of identifying genotyping diverse genomes of plant and animal systems including rice varieties, a kit thereof Patent: WO 03085133-A 36 16-OCT-2003; Centre for DNA Fingerprinting and Diagnostics, Centre for; the Department of Biotechnology, Ministry of Science & Technology Location/Qualifiers
                                                                                                                                                                                                                                                                                           synthetic construct synthetic construct
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Other publication FR 2749308 19971205
Location/Qualifiers
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NUCLEOTIDE PROBES AND METHOD FOR DETERMINING Patent: WO 9746700-A 31 11-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              unidentified unidentified
                                                                                                                                                                                                                                                 Nagaraju, J.G.
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Similarity 100.0%; Pred. No. 5.8e+04;
13; Conservative 0; Mismatches 0;
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   Conservative
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31 from Patent WO9746700.
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/mol_type="unassigned DNA"
/db_xref="taxon:32644"
                                                                   /organism="synthetic construct"
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/db xref="taxon:32630"
/note="A novel FISSR-PCR primer for
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100.0%; Pred. No. 5.7e+04;
vative 0; Mismatches 0;
4.1%; Score 13; DB 6; Length 18; 100.0%; Pred. No. 5.8e+04; tive 0; Mismatches 0; Indels
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BD103980/c
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       NISHIDA
OS Art
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PD 06-
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                                                                              Kit and method for determining HLA type
Patent: WO 0192572-A 84 06-DEC-2001;
NISSHINBO INDUSTRIES INC. SYSTEM RESEARCH INC, HIDETOSHI INOKO, TR
KAGIYA, TATSUO ICHIHARA, YOSHIYUKI MATSUMURA, SHOGO MORIYA, MICHIO
                                                                                                                                                                                                     BD103980
BD103980.1 GI:22649554
WO 0192572-A/84.
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                                                                                                                                     artificial sequences.
1 (bases 1 to 18)
Inoko,H., Kagiya,T., Ichihara,T., Matsumura,Y.,
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Patent: JP 2001321190-A 35 20-NOV-2001;
THE INSTITUTE OF PHYSICAL AND CHEMICAL RESEARCH,
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01-JUN-2001 WO 2001JF004662
01-JUN-2000 JF 00P 164798
HIDETOSHI INOKO,TAEKO KAGIYA,TATSUO ICHIHARA,YOSHIYUKI
                                        Artificial Sequence
WO 0192572-A/84
06-DEC-2001
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12-MAR-2001 JP 2001068285
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JP 2001321190-A/35
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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             RESULT 37
BD104190/c
  Locus
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KEYWORDS
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ACCESSION
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BD104081/c
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                                                                                                       Query Match
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BD104190
                                                                                                       l Similarity
13; Conserv
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PI SHOGO
PC C12Q1
CC Desci
FH Key
FT Sourc
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1 (bases 1 to 18)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              synthetic construct
synthetic construct
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WO 0192572-A/185.
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                                                                                CGCACGTCCTCCT 108
                                                          CGCACGTCCTCCT
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Justity 100.0%;
Conservative
                                                                                                       Conservative
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WO 0192572-A/185
06-DEC-2001
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Nishida, M.

Kit and method for determining HLA type
Patent: WO 0192572-A 185 06-DEC-2001;
NISSHINBO INDUSTRIES INC, SYSTEM RESEARCH INC, HIDETOSHI INOKO, TY
KAGIYA, TATSUO ICHIHARA, YOSHIYUKI MATSUMURA, SHOGO MORIYA, MICHIO
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PI SHOGO MORIYA, MICHIO NISHIDA

PC C1201/68, C12M1/00, C12N15/09, G01N33/53

PC CD Description of Artificial Sequence: capture

FH Key Location/Qualifiers

FT source 1.18
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C12Q1/68, C12M1/00, C12N15/09, G01N33/53
Description of Artificial Sequence: capture
Key Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-JUN-2001 WO 2001JP004662
01-JUN-2000 JP 00P 164798
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                                                                                                                                                                                                                                                                                                                       /organism='Artificial
                                                                                                                                                                                                                                                     organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Pred. No.
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Pred. No.
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5.8e+04;
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5.8e+04;
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PAT 27-AUG-2002
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AB068646/c
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JOURNAL
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JOURNAL
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Direct Submission
Submitted (04-AUG-2001) Akira Horii, Tohoku University Scho
Medicine, Molecular Pathology; 2-1 Seiryomachi, Aoba-ku, Sc
Miyagi 980-8575, Japan (E-mail:horii@mail.cc.tohoku.ac.jp,
Tel:81-22-717-8042, Fax:81-22-717-8047)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NISHIDA
OS Artif.
OS Artif.
PN WO 01:
PD 06-DE
PF 01-JUI
PR 01-JUI
PR 11DET
PI HIDET
PI SOGO
PC C1201.
CC Descript
FH Key
FT Source
FT Source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13,
                                                                                                                                                  A BAC-based STS-content map spanning a 35-Mb chromosome 1p35-p36 Genomics 74 (1), 55-70 (2001) 21269192
                                                                                                                                                                                                                Chen,Y.Z., Hayashi,Y., Wu,J.G., Takaoka,E., Maekawa,K., Watanabe,N., Inazawa,J., Hosoda,F., Arai,Y., Mizushima,H. Morohashi,A., Ohira,M., Nakagawara,A., Liu,S., Hoshi,M., and Soeda,E.
                                                                                                                                                                                                                                                                                             synthetic construct
synthetic construct
artificial sequence
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Synthetic construct DNA, at 1p36.
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Patent: WO 0192572-A 294 06-DEC-2001;
NISSHINBO INDUSTRIES INC. SYSTEM RESEARCH INC, HIDETOSHI INOKO, TE
KAGIYA, TATSUO ICHIHARA, YOSHIYUKI MATSUMURA, SHOGO MORIYA, MICHIO
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1 (Dases 1 to 18)

Inoko,H., Kagiya,T., Ichihara,T., Matsumura,Y., Moriya,S.
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synthetic construct
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WO 0192572-A/294.
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BD104190
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SHOGO MORIYA, MICHIO NISHIDA
C12Q1/68, C12M1/00, C12N15/09, G01N33/53
Description of Artificial Sequence: capture
Key Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-JUN-2001 WO 2001JP004662
01-JUN-2000 JP 00P 164798
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WO 0192572-A/294
06-DEC-2001
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/organism="synthetic construct"
/mol type="genomic DNA"
/db_xref="taxon:32630"
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Pred. No. 5.8
0; Mismatches
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                                                    Tohoku University School of eiryomachi, Aoba-ku, Sendai,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 18;
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    Query Match
Best Local
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AUTHORS
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AUTHORS
TITLE
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KEYWORDS
SOURCE
ORGANISM
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AR120194
LOCUS
DEFINITION
ACCESSION
VERSION
KEYWORDS
SOURCE
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DEFINITION
ACCESSION
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AR077249
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Best Local
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Livak, K.J. and (
Computer logic
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Sequence 7:
AR077249
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Livak, K.J. and Goodsaid, F.
Determination of a genotype of an amplification product
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13;
                                                                                                                                                                                                                        Sequence 7 from patent US AR120194
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unknown
                                                                                   Patent: US 6154707-A 7 28-NOV-2000;
Location/Qualifiers
                                                                                                                                                                        Unknown.
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Patent: US 5962233-A 7 05-OCT-1999;
                                                                                                                                                            Unclassified
                                                                                                                                                                                    Unknown.
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Similarity 100.0%;
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Similarity 100.0%; Pred. No.
13; Conservative 0; Mismatci
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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                                               /mol_type="unassigned
                                                             organism="unknown"
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Score 13;
Pred. No.
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Pred. No.
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red. No. 5.8e+04;
Mismatches 0;
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            DB 6;
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Similarity

100.0%;

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PAT 21-NOV-2001

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REFERENCE
AUTHORS
TITLE
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AR266141/c
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KEYWORDS
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ACCESSION
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VERSION
KEYWORDS
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E13797/c
LOCUS
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Matches 13
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                                                                                                                                                                                                                                                         140
                                                                                                                                                                                                                                                                                                                                                                                                                                                          strandedness: Single;
CC topology: Linear;
CC hypothetical: No;
CC anti-sense: Yes;
FH Key Loca
FT source
FT fource
                                                                                                                                                                                                                              16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           unclassified.

1 (bases 1 to 20)

1 (bases 1 to 20)

Ono,T., Mukaide,M., Hikichi,K. and Mizogami,M.

NEW OLIGONUCLEOTIDE, PRIMER FOR DISCRIMINATION IN GENOTYPE OF HEPATITIS C VIRUS COMPRISING THE SAME AND DISCRIMINATION IN GENOTYPE OF HEPATITIS C VIRUS BY USING THE PRIMER PATENT: JP 1997234072-A 49 09-SEP-1997;

S. R. L.KK.

S. R. L.KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OS Non
OC Art
PN JP
PD 09-
PF 01-
PR 01-
ONO TOMO
MASAFUMI
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1 (bases 1 to 20)
Bennett, C.F., Busch, H. and Wyatt, J.
Antisense modulation of GU protein expression
                                                                                                                         Sequence 40 from patent AR266141
                                                                                                                                                                                                                                                                                     l Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13;
                                               Unclassified
                                                              Unknown
                                                                              Unknown
                                                                                                         AR266141.1
                                                                                                                                                                                                                                                                                                                                                                                                                                   misc_feature
FT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primer for discriminating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  unidentified
                                                                                                                                                                                                                                                      CGAGGTTGCGACC 152
                                                                                                                                                                                                                              CGAGGTTGCGACC 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-SEP-1997
01-FEB-1996 JP 1996038875
01-FEB-1995 JP 95P 35997, 30-DEC-1995 JP 95P
TOMOYOSHI, MUKAIDE MASAKAZU, HIKICHI KAZUMASA, PI
                                                                                                                                                                                                                                                                                4.1%; Score 13; DB 6; L. larity 100.0%; Pred. No. 5.8e+04; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C12N15/09, C07H21/04, C12Q1/68, C12Q1/70, (C12N15/09, C12R1:92); CC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Artificial sequences. JP 1997234072-A/49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                 /note='Primer,OMM21TCS'
                                                                                                                                                                                                                                                                                                                                                        /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
                                                                                                         GI:29694987
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BD087778/c
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AX292891
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Best Local Similarity
Matches 13; Conserv
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OS Artif
PN JP 20
PD 20-NO
PF 12-MA
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                                                                                                                          A method of arraying genome clone
Patent: JP 2001321190-A 22 20-NOV-2001;
THE INSTITUTE OF PHYSICAL AND CHEMICAL RESEARCH, YUGENKAISHA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
AX292891
                                                                                                                                                                                       synthetic construct
synthetic construct
artificial sequences.
1 (bases 1 to 20)
                                                                                                                                                                                                                                                                                A method of arraying BD087778
                                                                                                                                                                                                                                                  JP 2001321190-A/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Barany, F., Zirvi, M., Gerry, N.P., Favis, R. an Method of designing addressable array for de sequence differences using ligase detection Patent: WO 0179548-A 4653 25-CCT-2001; CORNELL RESEARCH FOUNDATION, INC. (US)
                                                                                                                                                                                                                                                               BD087778.1 GI:22633388
                                                                                                                                                                                                                                                                                                             BD087778
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Location/Qualifiers
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C12N15/00,
C12N15/00
                                                                                                                                                                                                                                                                                                                                                                                  GAGTGGACCATCG 16
                                                                    Artificial Sequence
JP 2001321190-A/22
20-NOV-2001
                          C12N15/09, C12N15/09, C12M1/00, C12Q1/68, G01N33/53, G01N33/566,
                                          12-MAR-2001 JP 2001068285
EIICHI SOEDA
                                                                                                                                                                                                                                                                                                                                                                                                                                        4.1%; Score 13; DB ilarity 100.0%; Pred. No. 5. Conservative 0; Mismatches
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4653 from Patent WO0179548.
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  note="Hypothetical Probe Sequence"
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5.8e+04;
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PAT 27-AUG-2002

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RESULT 46
E13781/c
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Best Local S
Matches 13
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Submitted (04-AUG-2001) Akira Horii, Tohoku University Sch
Medicine, Molecular Pathology; 2-1 Seiryomachi, Aoba-ku, S
Miyagi 980-8575, Japan (B-mail:horii@mail.cc.tohoku.ac.jp,
Tel:81-22-717-8042, Fax:81-22-717-8047)
 E13781 21 bp DNA PCR primer for discriminating genotype la
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       and Soeda,E.

A BAC-based STS-content map spanning a 35-Mb region of human chromosome 1p35-p36

Genomics 74 (1), 55-70 (2001)

21269192
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                                                                                                                            GACTGGGGATGTC 316
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (bases 1 to 20)
                                                                                                                                                         4.1%; Score 13; DB 12; 
larity 100.0%; Pred. No. 5.8e+04;
Conservative 0; Mismatches 0;
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Conservative (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bource
                                                                                                                                                                                                                                   /note="reverse primer for human STS sts-WI-13136 at 1p36 sts-WI-13136 obtained from clones B72G3, B37DI16, B34IF17 B26SD10, B34IF17, B53F6, B128O2, B85K5, B73C3, Human BAC library RPCI-11"
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/mol_type="genomic DNA"

/db_xref="taxon:32630"
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                                                                                                                                                                                                                                                                                                                  xref="taxon:32630"
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linear PAT 27-1
of HCV (Hepatitis
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STS sts-WI-13136
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E13781
E13781.1
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Ono,T., Mukaide,M., Hikichi,K. and Mizogami,M.
Ono,T., Mukaide,M., Hikichi,K. and Mizogami,M.
NEW OLIGONUCLEOTIDE, PRIMER FOR DISCRIMINATION IN GENOTYPE OF HEPATITIS C VIRUS THE SAME AND DISCRIMINATION IN GENOTYPE OF HEPATITIS C VIRUS BY USING THE PRIMER PATENT: JP 1997234072-A 33 09-SEP-1997;
                                                                                                                                                       Ono,T., Mukaide,M., Hikichi,K. and Mizogami,M.
NEW OLIGONUCLEOTIDE, PRIMER FOR DISCRIMINATION IN GENOTYPE
HEPATITIS C VIRUS COMPRISING THE SAME AND DISCRIMINATION IN
GENOTYPE OF HEPATITIS C VIRUS BY USING THE PRIMER
Patent: JP 1997234072-A 45 09-SEP-1997;
                                                                                                                                                                                                                                                                                                             PCR primer for discriminating E13793 .1 GI:3252561
                                                  PR PR OC OS
                                                                                                                                          S R L:KK
                                                                                                                                                                                                                                                                                                  E13793.1 GI:3252561 JP 1997234072-A/45.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PF PO OS
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                        Artificial sequences.

JP 1997234072-A/45
09-SEP-1997
01-FEB-1996 JP 1996038875
01-FEB-1995 JP 95P
01-FEB-1995 JP 95P
TOMOYOSHI, MUKAIDE MASAKAZU, HIKICHI KAZUMASA, PI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Artificial sequences. JP 1997234072-A/33 09-SEP-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Key
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topology: Linear;
hypothetical: No;
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01-FEB-1995 JP 95P 359
 C12N15/09, C07H21/04, C12Q1/68, C12Q1/70, (C12N15/09, C12R1:92);
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100.0%; Pred. No. 5.9e+04;
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PAT 27-APR-1998

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                              Greaves,D., Price,S. and Watkins,H. Functional genetic variants Patent: WO 0246462-A 8 13-JUN-2002; Isis Innovation Limited (GB)
                                                                                                                                                                       AX458691 21 bp
Sequence 8 from Patent WO0246462.
AX458691
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Sequence 2
AX015616
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/mol_type="unassigned DNA"
/db xrefs"ttaxon:32630"
/noTe="synthetic oligonucleotid
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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AR236424
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Unclassified.

1 (bases 1 to 22)

1 (bases 1 to 22)

Goldfine, I., Trischitta, V., Vigneri, R., Pizzuti, A. and Prittitta, L.A.

Polymorphic human PC-1 sequences associated with insulin Patent: US 6465185-A 10 15-OCT-2002;
                                                                                                                                                AR236424
Sequence 10
AR236424
                                                                                                                                                                                                                                                                                                                                                                                                                                     PF 25-AUG-2000 WO 2000JP005757
PR 05-OCT-1999 JF 99P 2844001
PFI YASUHIRO FURUICHI, AKERA SHIMAMOTO, SAORI KITAO, KAORI NISHIKAWA
DEGC C12N15/12, C12N9/14, C12Q1/68, C07K16/18, G01N33/53, A01K67/00 CC
DESCRIPTION OF Artificial Sequence; Artificially Synthesized CC
Primer Sequence
FH Key
                                                                                           Unknown
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1 (bases 1 to 21)

Furuichi,Y., Shimamoto,A., Kitao,S. and Nishikawa, Nucleus localizing RecQ5-type DNA helicase Patent: WO 0125425-A 45 12-APR-2001;
AGENE RESEARCH INSTITUTE CO LTD, YASUHIRO FURUICHI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    82 GGAGGAGGGCGAG 94
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WO 0125425-A/45.
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WO 0125425-A/45
12-APR-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers.
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                                                                                                                                   GI:27280403
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REFERENCE AUTHORS TITLE JOURNAL

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KEYWORDS SOURCE DEFINITION ACCESSION VERSION

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                                                                                                                                            Patent:
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Resnick,R.M. and Youn
Methods, primers and
variants
                                                                                                                                                                                                                                 Sequence 21 from patent I22162 I22162.1 GI:1602516
                                                                                                                                                                                                                                                                                                                                                                                                                                      1 (bases 1 to 23)
Chenchik, A., Jokhadze, G. and Bibilashvilli, R.
Methods of assaying differential expression
Patent: US 5994076-A 773 30-NOV-1999;
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Sequence 773 from patent
AR090653
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Similarity 100.0%; Pred. No.
13; Conservative 0; Mismatci
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ilarity 100.0%;
Conservative (
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llarity 100.0%;
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Location/Qualifiers
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/mol_type="unassigned DNA"
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/mol_type="genomic"
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                                                                                                                                                              Young, K.K.Y. and probes for detection
                                                       score 13; DB
f; Pred. No. 5.9
f; Mismatches
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Pred. No. 5.9e+04;
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AR259842
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62 CGGCTCCTGCGTC
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Chenchik,A., Jokhadze,G. and Bibilashvilli,R.
Methods of assaying differential expression
Patent: US 6489455-A 773 03-DEC-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AR259842
Sequence 773 from patent
AR259842
                      1 (bases 1 to 24)
Resnick, R.M. and You
Methods, primers and
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Chenchik, A., Jokhadze, G. and Bibilashvilli, R.
Methods of assaying differential expression
Patent: US 6352829-A 773 05-MAR-2002;
Patent: US 5527669-A 14 18-JUN-1996;
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Similarity 100.08;
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/mol_type="genomic |
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                        Young, K.K.Y. and probes f
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AX288258
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                                                                                                            Ulfendahl, P.J. and Wong, K.C. Primers for identifying typing or classifying Patent: WO 0065088-A 64 02-NOV-2000; Amersham Pharmacia Biotech AB (SE) Location/Qualifiers
                                                                                                                                                                                                         synthetic construct synthetic construct
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Sequence 64 from Patent
AX042498
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Sequence 20 from Patent
AX288258
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                                                 /db_xref="t/note="DBQ
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/db_xref="teaxon.32630"
/noTe="Hypothetical Probe Sequence"
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/mol_type="unassigned DNA"
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                                              heterozygote typing
Score 13;
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AX043378/c
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artificial sequences.
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                                                                                                                                                                              pathogenic fungi
Patent: EP 1118677-A 3
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AX043378
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Primers for identifying typing or classifying nucleic Patent: WO 0065088-A 944 02-NOV-2000;
Amersham Pharmacia Biotech AB (SE)
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                                                                                                 /organism="synthetic construct"
/mol_tyse="unassigned DNA"
/mol_tyse="taxon:32630"
/noTe="Synthetic Oligonuclectid
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/db xref="taxon:32630"
/note="DQB1 Heterozygote Primer
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from Patent EP1118677.
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Pred. No. 6e+04;
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AX354462/c
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              Homo sapiens (human)
Homo sapiens
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AX354462
AX354462.1 GI:18619304
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Sequence 3 from Patent WO0153521
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 Eukaryota; Metazoa; Chordata; Craniata;
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/mol_type="unassigned DNA"
/mol type="unassigned DNA"
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/db xref="taxon:32630"
/note="Synthetic Oligonucleotide"
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Sequence 2209 from Patent
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Sequence 2208 from Patent
AX500901
                                                Human testis expressed patched like protein Patent: EP 1229046-A 2209 07-AUG-2002;
                                                                                                                   Eukaryota;
                                                                                                                              Homo sapiens
                                                                                                                                        Homo sapiens (human)
                                                                                                                                                                  AX500902.1 GI:23383195
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                                                                           Zhan, J
                                                                                                    Eukaryola; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Similarity 100.0%; F
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/organism="Homo sapiens"
                        Location/Qualifiers
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Primates;
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Local Similarity 100.0%; Pred. No. 6e+04;
hes 13; Conservative 0; Mismatches
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AX500904
AX500904.1 GI:23383197
                                                                                                                                                                      Human testis expressed patched like protein Patent: EP 1229046-A 2211 07-AUG-2002;
                                                                                                                                                                                                                  Homo sapiens (human)
Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 2210 from Patent
AX500903
AX500903.1 GI:23383196
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         GGAAGGCAAGCAG 37
                              4.1%; So larity 100.0%; I Conservative 0;
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/mol_type="unassigned DNA"
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                             Score 13; DB 6; ; Pred. No. 6e+04; 0; Mismatches
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Mammalia; Eutheria; Primates;
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Sequence 2212 from Pater
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Human testis expressed patched like protein Patent: EP 1229046-A 2214 07-AUG-2002; Aeomica, Inc. (US)
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Sequence 2216 from Patent
AX500909
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Mammalia; Eutheria;
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Sequence 2217 from Patent
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Mammalia; Eutheria;
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Similarity 100.0%; Pred. No. 6e+04;
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Search completed: February 2, 2005, 20:25:58 Job time: 1357.61 secs
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Sequence 2219 from Patent EP1229046.
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AX500912.1 GI:23383205
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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ADB22460
ABB28618
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ADK74647
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Adk01333 Rat DNA m
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Abd24848 Al092623-
Adh67348 Human glu
Adh67400 Human glu
Adh67400 Human glu
Adh74647 Chimeric
Adm14246 Human mpg
Adm99302 Stem cell
Aaq75649 Reverse t
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Aax84758 PCR prime
Aad34264 Human CYP
Aad26900 Bacterial
Abk86170 Oligo dT
Aax78723 Human pan
Aad26899 Bacterial
Aax78723 Human pan
Aad26899 Bacterial
Aad39650 PolypNP
Aax71936 Anchored
Aah41080 Nucleotid
Abd79879 Nucleotid
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Mammalian
52 Stem cell
51 Universal
11 Human SCF
51 Mammalian
72 Human SCF
7 Human SCF
7 Human SCF
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18 SCF unive
8 SCF unive
18 SCF unive
18 Human cli
18 Human cli
48 A1092623
48 Human glu
00 Human glu
01 Human glu
01 Human glu
01 Human glu
02 Chimeric
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JP06303997-A

Synthetic

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RESULT 1
AAQ75652/c
ID AAQ75652 standard; DI
XX
AC AAQ75652;
XX
DT 04-AUG-1995 (first of the text)
XX
COT 04-AUG-1995 (first of the text)
XX
Analysis; gene expresion
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Analysis; restriction
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105
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107
Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
                           Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                               (first entry)
                                                                                    DNA;
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                                                                                     ВP
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ADB37210
ADB37236
ADE77617
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AAF6020
AAL45359
AAL763378
AAR76221
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AAR76222
AAR7669
AAQ7569
AAX199469
AAX1994
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ABL39401
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ABS53437
ABA93239
AAD56446
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Adi34189 Nucleotid
Adh78590 Test elem
Ado28710 Single st
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ARESULT 2
ADX013/C
ID ADX013
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AC ADX013
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DT 06-MAY
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B8; hy
KW B8; hy
KW blood;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stransed cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                            Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.
                                                                                                                                                                                                                                                                                                                                                                                                                      04-SEP-2003.
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                                                                                                                                                                                             Boekenkamp D,
                                                                                                                                                                                                                                                                                                         28-FEB-2002; 2002DE-01008794.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Analysis of cDNA and gene expression -
by digestion with restriction enzymes.
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cell; food additive; food supplement.
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Example;

Page

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gpp;

German.

describes a novel method for sorting single-stranded

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AAQ75570/c
                                                                                                                                                                                                                                                                     Query Match
Best Local
                                                                                                                                                                                                                                                               Matches
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                                                                                             JP06303997-A
                                                                                                                                                                                            AAQ75570 standard; DNA; 20
                                                                                                                                                                                                                                                                                            Sequence
Analysis of cDNA and gene expression - by digestion with restriction enzymes.
                     WPI; 1995-018287/03
                                                                 16-APR-1993;
                                                                                                            Synthetic
                                                                                                                         aggregate;
                                                                                                                          Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
                                                                                                                                                Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                               04-AUG-1995
                                                   16-APR-1993;
                                    ) NIPPON
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                                                                                                                                                                                                                                                                                                          probes used in the method of the invention.
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                                                                                                                                                                                                                                                                                            21 BP; 0
                                                                                                                                                                                                                                                               Conservative
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                                                   93JP-00112515
                                    TELEGRAPH & TELEPHONE CORP
                                                                 93JP-00112515
                                                                                                                                                                                                                                                                                            A; 1 C; 1 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                     100.0%;
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                                                                                                                                                                                                                                                                      Score 21;
Pred. No.
                                                                                                                                                                                                                                                              Mismatches
      by amplification of mRNA followed
                                                                                                                                                                                                                                                                    DB 10; Length 21; 1.1e+03;
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The present sequence is that of a 5' PCR primer, used with the 3' primer given in AAF30362, to amplify a cDNA fragment corresponding to nucleotides 904-172 of the human chfr sequence given in AAF30352. The chfr gene encodes the human mitotic checkpoint protein Chfr (see AAB20219), which is required for regulation of the transition of Cells from prophase to metaphase during mitosis. Loss of expression of Chfr is associated with a predisposition to tumourigenesis upon exposure to mitotic stress. A set of primers (see AAF30353-76) was used to amplify regions spanning the entire chfr coding region in order to determine whether the chfr gene is mutated in any of the human cancer cell lines \$M480, DLD1, HT29, HCT116, \$ASOS2, UZOS, IMFS and NGP. A mutation leading

Novel nucleic acid sequence of mitotic checkpoint gene encoding a checkpoint with forkhead-associated domain and ring finger protein, diagnosing tumorigenic cells and in screening for anticancer drugs.

for

Example 3; Page 38; 85pp; English.

WPI;

2001-182927/18

Halazonetis T,

Scolnick

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WISTAR

INST ANATOMY & BIOLOGY:

to a Val-580 to Met amino acid substitution was identified in the chfr gene of U2OS cells. Chfr polypeptides and chfr nucleic acids are used

ed in which

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CC nucleic acids by isolation and hybridisation of nucleic acid pools, then comprise variable and non-variable regions. The capture oligonucleotides agents that are (a) immobilised on the surface of a solid matrix and (b) CC comprise variable and non-variable regions. The capture oligonucleotides comprises a sirable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3 'variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow that comprises a sequence of 10-50, particularly 15-25, T residues. The capture CC agents are particularly locked nucleic acids (LNA) and the anchor region CC comprises a sequence of 10-50, particularly 15-25, T residues. The CC capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, CC interaction with streptavidin. The matrix is of plastic, ceramic, glass, CC interaction with streptavidin material and/or membrane, having semical particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, CC is particularly a layer of (bio)molecular filaments and binding of single extranded nucleic acids to the surface is (quasi)covalent, supramolecular, CC is partity of pharmaceuticals and (i) for analysis of patterns, especially in mucosal, CC hair root, blood, nerve or germ cells and (ii) for determining the CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and call ows imaging of the complete sapid, inexpensive and reproducible call was imaging of the comparison of nucleic acid sfrom cells. It CC allows imaging the particular probable and comparison of nucleic acid pool. Since the CC are the comparison of such the nucleic acid pool. Since the CC are the comparison of such the nucleic acid pool. Since the comparison is possible. ADX01281-ADX01344 represent
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AAF30361
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RESULT 5
AAF30363
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                                                                                   CC The present sequence is that of a 5' PCR primer, used with the 3' primer CC given in AAF30364, to amplify a cDNA fragment corresponding to CC nucleotides 904-1902 of the human chfr sequence given in AAF30352. The CC chfr gene encodes the human mitotic checkpoint protein Chfr (see CC AAB20219), which is required for regulation of the transition of Chfr is CC remover the control of the constitution of Chfr is CC associated with a predisposition to tumourigenesis upon exposure to CC mitotic stress. A set of primers (see AAF3033-76) was used to amplify CC regions spanning the entire chfr coding region in order to determine CC whether the chfr gene is mutated in any of the human cancer cell lines CC waf80, DLD1, HT29, HCT116, SAOS2, U2OS, IMR5 and NGP. A mutation leading CC to a Val-580 to Met amino acid substitution was identified in the chfr Gene of U2OS cells. Chfr polypeptides and chfr nucleic acids are used in CC methods of diagnosing tumourigenic cells and of screening for drugs which CC can inhibit the activity of Chfr in a cancer cell, rendering it more CC sensitive to additional antitumour therapies
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Pred. No.
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Conservative

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Mismatches

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Gaps

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501 CAAGGAAAATGTGTTCCATG

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The present sequence is that of a 3' PCR primer, used with the 5' primer CC given in AAF30353, to amplify a cDNA fragment corresponding to nucleotides 66-562 of the human entry sequence given in AAF30352. The chfr Cg gene encodes the human mitotic checkpoint protein Chfr (see AAB20219), CC which is required for regulation of the transition of cells from prophase to metaphase during mitosis. Loss of expression of Chfr is associated CC with a predisposition to tumourigenesis upon exposure to mitotic stress. CA set of primers (see AAF30353.76) was used to amplify regions spanning the entire chfr coding region in order to determine whether the chfr gene is mutated in any of the human cancer cell lines Sw440, DLDI, HT29, CC HCT116, SAOS2, U2OS, IMRS and NGP. A mutation leading to a Val-580 to Met CC chfr polypeptides and chfr nucleic acids are used in methods of CC chfrosting tumourigenic cells and of screening for drugs which can consisting tumourigenic cells and of screening for drugs which can cancer cell, rendering it more
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                                                 20 BP; 5
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RESULT 8
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                Analysis; gene 'expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (WIST-) WISTAR INST ANATOMY & BIOLOGY.
                                                                                                                                                                                                                                                                                                862 AAGAAAATGAGAGGAGATGG 881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       checkpoint gene chfr 5'
                                                                                                                                                                                                                                                                 ш
                                                                     transcription primer used
                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                    20 BP; 10 A; 0 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                AAGAAAATGAGAGGAGATGG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                         it the activity of Chfr in a cancer cell, to additional antitumour therapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tumorigenic cells and in screening
                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                     100.0%;
                                                                                                                                                                                                                                                                                                                                                                     0.7%;
                                                                                                                                                                            21
                                                                                                                                                                            ВÞ
                                                                                                                                                                                                                                                                                                                                   0,
                                                                                                                                                                                                                                                                                                                                                   Score 20; Pred. No.
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                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                 in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     primer
                                                                                                                                                                                                                                                                                                                          DB 5; 4...
0. 2.5e+03;
0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          for anticancer drugs.
                                                                                                                                                                                                                                                                                                                                                                    Length 20
                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            finger; Chfr; human;
drug screening;
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                                                                                                                                                                                                                                                                                                                                 Gaps
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RESULT 9
AAQ75653/c
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ filles AAQ755978) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                               Analysis; ; aggregate;
                                                             Analysis of
                                                                                                                         16-APR-1993;
                                                                                                                                                                   01-NOV-1994.
                                                                                                                                                                                                                                                                                  04-AUG-1995
                                                                                                                                                                                                                                                                                                       AAQ75653;
                                                                                                                                                                                                                                                                                                                           AAQ75653 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 0 A; 1 C; 2 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Analysis of cDNA and gene expression - by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic: ;
                                                                                 WPI; 1995-018287/03.
                                                                                                      (NITE )
                                                                                                                                             16-APR-1993;
                                                                                                                                                                                       JP06303997-A.
                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                              Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 6; 11pp; Japanese
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-APR-1993;
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                                                  digestion
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                                                                                                                                                                                                                               gene expression; reverse
; restriction enzyme; ss.
                            Page
                                                             cDNA and
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                                                  with restriction
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                                                                                                     TELEGRAPH
                                                                                                                          93JP-00112515
                                                                                                                                              93JP-00112515
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                            6; 11pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                0.7%;
                                                                                                                                                                                                                                                                                                                            21
                                                                                                     & TELEPHONE
                             Japanese
                                                             expression -
                                                                                                                                                                                                                                                                                                                            ВP
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Pred. No.
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                                                                                                                                                                                                                                         transcription; primer; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                  DB 2; L.
5. 2.5e+03;
0;
                                                          λq
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                                                           amplification
                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
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                                                           mRNA followed
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A method for the analysis of cDNA comprises (a) double-stranded cDNAs by using an aggregate of

mRNAs and a plural type

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AAQ75654/c
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Best Local
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Matches 20
                                                    Matches
                                                                                                                       A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type clabelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                      Sequence
                                                                                                                                                                                                                                                  Disclosure; Page 6; llpp; Japanese.
                                                                                                                                                                                                                                                                                    Analysis of cDNA and gene expression - by amplification of mRNA followed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                       WPI; 1995-018287/03.
                                                                                                                                                                                                                                                                                                                                                                        16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                    16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                              01-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                       JP06303997-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Analysis; gene expression; reverse transcription; primer; cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                            (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         aggregate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAQ75654;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ75654 standard;
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                                                                 Local
                                                                                                                                                                                                                                                                                digestion with
                           2659
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20; Conser
                       GACAAAAAAAAAAAAAAA 2678
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GACAAAAAAAAAAAAAAAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       restriction enzyme; ss.
                                                                                                      BP; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                         93JP-00112515.
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                                                                                                      P,
                                                                                                                                                                                                                                                                               restriction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
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100.0%; Pr
                                                              0.7%;
                                                                                                  2 C; 1 G; 18 T; 0 U; 0 Other;
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Pred. No.
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                                                   Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 U; 0 Other;
                                                                            DB 2;
                                                                2.5e+03
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                                                                         Length 21
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                                               Gaps
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RESULT 11

B ADK01309/c

ADK01309

standard;

DNA;

21

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Best Loc Matches Query Match

Local

Similarity

0.7%;

Score 20; Pred. No. Mismatches

DB 10; 2.5e+03;

Length 21; Indels

<u>.</u>

Gaps

0

0;

АСААЛААЛААЛААЛААЛ 2679

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Comprise variable and non-variable region. The complement of which is present at CC have a 5'-invariable anchor region, the complement of which is present at CC that comprises all possible combinations of up to 10 nucleotides to allow CC binding of particular sorts of single stranded nucleic acids. The capture CC agents are particularly locked nucleic acids (LNA) and the anchor region CC capture oligonucleotides are biotinnylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, CC metal, resin, gel, crystalline material and/or membrane, having semi-CC conducting properties and especially in the form of a chip. Its surface by interaction with streptavidin. The material and/or membrane, having semi-CC conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single crystacl, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the captures. The method provides rapid, inexpensive and reproducible acids (amino, carboxylic or fatty acid) or their derivatives, salts and captures. The method provides rapid, inexpensive and reproducible captures in again of the complete pattern of all nucleic acids from cells. It captures is based on comparison of nucleic acid pools, not individual represent the method is based on comparison is possible. ADK01281-ADK01344 represent components and nucleic acid sold in a cell, and capture probes used in the method of the invention.
        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example; Page 5; 8pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and constant regions.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          patterns and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Boekenkamp
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-SEP-2003.
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blood; nerve; germ cell; food additive; food
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rat DNA microarray capture oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              single-stranded nucleic acid, and screening active agents,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DEGUSSA BIOACTIVES GMBH.
            21
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        BP; 1
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        Ą
0 C; 1
    G; 19
T; 0
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U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          supplement
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RESULT 12 ADK01312/c

ADK01312 standard; DNA; 21 BP.

06-MAY-2004 ADK01312;

(first entry)

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X5XFXBXFFX8XFXFXFXFXFXFXFFFX8X9S9S9S9S9S9S9S9S9S9S9S9S9S
                                                                                                                                         This invention describes a novel method for sorting single-stranded conucleic acids by isolation and hybridisation of nucleic acid pools, then care calding out, where the nucleic acids are selectively bound using capture comparise that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides comprise variable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region comprises once in each nucleic acid and a 3'-variable, discriminatory region comprises a sequence of 10-50, particularly 15-25, T residues. The capture capture oligonucleotides to allow binding of particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, complete acids to the surface is (quasi) covalent, suprample collar, ghysical, resin, gel, crystalline material and/or membrane, having semi-conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramplecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (anion, carboxylic or fatty acid) or their derivatives, salts and conjusted in the mothod of the invention. Since the capture probes used on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADKO1381-ADKO1344 represent comparison of nucleic acid pools, and individual in the method of t
        Best Loc
Matches
                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28-FEB-2002; 2002DE-01008794
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                            Local Similarity
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100.0%; --
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                                                                                                                                    method
                                               Score 20;
                              Pred. No.
                                                                                         20 T; 0
                                                                                                                                  the invention.
                                                                                         U; 0 Other;
                       2.5e+03;
                                               DB 10; Length 21;
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Conservative

Mismatches

Indels

0

Gaps

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Sequence 21

BP; 0

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                             Comprise variable and non-variable regions. The capture oligonucleotides comprise variable and non-variable regions. The capture oligonucleotides comprise variable and non-variable regions. The capture oligonucleotides comprise variable anchor region, the complement of which is present at compare once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture comprises a sequence of 10-50, particularly 15-25, T residues. The capture comprises a sequence of 10-50, particularly 15-25, T residues. The capture comprises a sequence of 10-50, particularly 15-25, T residues. The capture comprises a sequence of 10-50, particularly 15-25, T residues. The capture comprises a sequence of 10-50, particularly 15-25, T residues. The computed on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, comparticularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, comparticularly an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, cardivity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and cardivity of patterns of a comparison of a comparison of all nucleic acids from cells. It can detect very small differences in the nucleic acids from cells, and can detect very small differences in the nucleic acid pool. Since the comparison of nucleic acid pools, not individual capture probes used in the method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example; Page 5; 8pp; German
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cc nucleic acids by isolation and hybridisation of nucleic acid pools, then comprise that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides comprise variable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow that comprises a sequence of 10-50, particularly 15-25, T residues. The capture comprises a sequence of 10-50, particularly 15-25, T residues. The capture comprises a sequence of 10-50, particularly 15-25, T residues. The capture comprises a sequence of 10-50, particularly 15-25, T residues. The capture comprises a sequence of 10-50, particularly 15-25, T residues. The capture comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, aciditives or supplements, especially minerals, trace elements, organic aciditives or supplements, especially minerals, trace elements, organic aciditives in aciditives, salts and compunity of pharmaceuticals and/or nutritional compounds, e.g. food cativity of pharmaceuticals and/or nutritional compounds. E.g. food cativity of physical cativity of physical cativity of p
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2.5e+03;
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310 CTGGAAGATACCAGCACCAG 329

Matches Query Match Best Local

Local

Similarity

0.7%;

DB 5; 1

Length 28

Indels

0

Gaps

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0; Mismatches Score 20; Pred. No. Sequence

28

BP; 12 A; 6 C; 6 G; 4 T; 0 U; 0 Other;

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RESULT 15
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                  chir gene encodes the human mitotic checkpoint protein Chfr (see AAB20219), which is required for regulation of the transition of cells from prophase to metaphase during mitosis. Loss of expression of Chfr is associated with a predisposition to tumourigenesis upon exposure to mitotic stress. A set of primers (see AAF30333-76) was used to amplify regions spanning the entire chfr coding region in order to determine whether the chfr gene is mutated in any of the human cancer cell lines SW480, DLD1, HT29, HCT116, SAOS2, UZOS, IMR5 and NGP. A mutation leading to a Val-580 to Met amino acid substitution was identified in the chfr gene of UZOS cells. Chfr polypeptides and chfr mucleic acids are used in methods of diagnosing tumourigenic cells and of screening for drugs which can inhibit the activity of Chfr in a cancer cell, rendering it more sensitive to additional antitumour therapies
                                                                                                                                                                                                                                                                                                                               Novel nucleic acid sequence of mitotic checkpoint gene encoding a checkpoint with forkhead-associated domain and ring finger protein, diagnosing tumorigenic cells and in screening for anticancer drugs.
                                                                                                                                                                                                                                   nucleotides 352-1055 of the human chfr sequence given in AAF30352.
                                                                                                                                                                                                                                                   The present sequence is that of a 5' PCR primer, used with tgiven in AAF30356, to amplify a cDNA fragment corresponding
                                                                                                                                                                                                                                                                                                        Example 3; Page 38; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-182927/18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                primer.
                                                                                                                                                                                                                                                                    PCR primer, used with the 3' primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.5e+03
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RESULT 16
AAF30356/c
RESULT 17
AAQ83940
ID AAQ83
XX
AC AAQ83
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                                                                                                                                                                       Query Match
Best Local S
Matches 20
                                                                                                                                                                                                                                                              associated with a predisposition to tumourigenesis upon exposure to mitotic stress. A set of primers (see AAF30353-76) was used to amplify regions spanning the entire cofir coding region in order to determine whether the chfr gene is mutated in any of the human cancer cell lines SW480, DLD1, HT29, HCT116, SAOS2, U2OS, IMR5 and NGP. A mutation leading to a Val-580 to Met amino acid substitution was identified in the chfr gene of U2OS cells. Chfr polypeptides and chfr nucleic acids are used in methods of diagnosing tumourigenic cells and of screening for drugs which can inhibit the activity of Chfr in a cancer cell, rendering it more sensitive to additional antitumour therapies
                                                                                                                                                                                                                                                                                                                                                                                                                         given in AAR30355, to amplify a cDNA fragment corresponding to nucleotides 352-1055 of the human chfr sequence given in AAR30352. The chfr gene encodes the human mitotic checkpoint protein Chfr (see AAB20219), which is required for regulation of the transition of cells from prophase to metaphase during mitosis. Loss of expression of Chfr is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel nucleic acid sequence of mitotic checkpoint gene encoding a checkpoint with forkhead-associated domain and ring finger protein, diagnosing tumorigenic cells and in screening for anticancer drugs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Checkpoint with forkhead associated domain and ring finger; Chfr; human; mitosis; cell cycle; tumour; diagnosis; antitumour; drug screening; ubiquitin-protein ligase; PCR primer; ss.
 AAQ83940;
                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence is that of a 3' given in AAF30355, to amplify a cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Halazonetis T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-MAY-2001
                               AAQ83940 standard; DNA; 30 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 3; Page 38; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-JUN-2000; 2000WO-US016391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200109150-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human checkpoint gene chfr 3' PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAF30356;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAF30356
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (WIST-) WISTAR INST ANATOMY & BIOLOGY.
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                                                                                                                                                                                        Similarity
                                                                                                                                          CTGACATGCATCATCTGCCA 1013
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         standard;
                                                                                                                                                                                                                                      28 BP; 10
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                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Scolnick D
                                                                                                                                                                                                                                    A; 4 C; 8 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
                                                                                                                                                                                      0.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hat of a 3' PCR primer, used with the lify a cDNA fragment corresponding to
                                                                                                                                                                       0,
                                                                                                                                                                                      Score 20;
Pred. No.
                                                                                                              ø
                                                                                                                                                                       Mismatches
                                                                                                                                                                                      DB 5; L, 2.4e+03;
                                                                                                                                                                       0
                                                                                                                                                                                                     Length 28
                                                                                                                                                                       Indels
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                                                                                                                                                                       Gaps
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RESULT 18
AAF60462
ID AAF60
XX Oliga
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                                                                                                                                                                                                                                                                                                                                   The sequences given in AAQ83938, AAQ83952 and AAQ83940 are used in the CC construction of an oligonucleotide clamp. The clamp is a comb-type CC branched polymer which has 3' termini and was used to bind a target CC sequence comprising a segment of the HIV pol and nef genes in single stranded or double stranded forms. An oligonucleotide clamp is a compound capable of forming a covalently closed macromolecule or a stable circular complex after specifically binding to the target polymucleotide. Oligonucleotide clamps generally comprise one or more oligonucleotide moieties capable of specific binding to the target molecule and one or conceptairs of binding moieties covalently linked to the oligonucleotide moieties. Upon annealing of the oligonucleotides moieties to the target plymucleotide, the binding moieties of a pair are bought into correct pairs of the interaction of the binding moieties effectively clamps the specifically annealed oligonucleotide moieties to the target the specifically annealed oligonucleotide moieties to the target polymucleotide. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                Query Match
Best Local 8
                             Oligonucleotide clamp #22.
                                                              27-APR-2001
                                                                                           AAF60462
                                                                                                                           AAF60462 standard; DNA; 30
                                                                                                                                                                                                                                                                                                                                 Sequence 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 8; Page 33; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthesis of branched polymers and novel branched polymeric structures used as molecular probes esp. for detecting poly-nucleotide(s).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1995-060944/08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-JUL-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HIV; pol; nef; oligonucleotide clamp; branched; macromolecule; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oligonucleotide clamp o,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-MAR-2003.
04-OCT-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (LYNX-) LYNX THERAPEUTICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12-JAN-1995.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9501365-A1
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                                                                                                                                                                                                                                   2660 АСААЛАЛАЛАЛАЛАЛАЛА 2679
                                                                                                                                                                                                        4
                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                        ACAAAAAAAAAAAAAAAAA 23
                                                                                                                                                                                                                                                                                                                                 BP;
                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (revised)
                                                            (first entry)
                                                                                                                                                                                                                                                                                                                              27 A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /*tag=  a
/note= "Modified with SP(O-) (=O)-"
                                                                                                                                                                                                                                                                                 0.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   entry)
                                                                                                                                                                                                                                                                                                                               3 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for producing comb-type brached
                                                                                                                           ВP
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                                                                                                                                                                                                                                                                                 Score 20; DB 2;
Pred. No. 2.4e+(
                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                   2.4e+03;
                                                                                                                                                                                                                                                                                                Length 30;
                                                                                                                                                                                                                                                                   Indels
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                                                                                                                                                                                                                                                                   Gaps
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Oligonucleotide clamp; ds

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RESULT 19
AAQ75549/c
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                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Analysis of cDNA and
                        WPI; 1995-018287/03.
                                                                         16-APR-1993;
                                                                                                16-APR-1993;
                                                                                                                         01-NOV-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to a method for synthesising a branched or multiply connected macromolecular structure, comprising oligonucleotide clamps (OC). The macromolecular structure is capable of specifically binding to a target molecule, and can therefore be used as probes. At least one OC comprises a target binding sequence that binds specifically and stably with the target molecule, and at least two OCs comprise signal generation moieties capable of generating a detectable signal in the presence of the target molecule. In addition the OCs are connected to one another by thioalkylamino, or thiophosphorylalkylamino bridges. The present sequence is an OC used in the present invention
                                                                                                                                                  JP06303997-A
                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                         Analysis; gene expression; reverse transcription;
                                                                                                                                                                                                                                                               04-AUG-1995
                                                                                                                                                                                                                                                                                                              AAQ75549 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthesizing branched nucleic acids useful as diagnostic and molecular probes, involves combining first units having haloalkylamino groups and second units having thiol or phosphorothicate groups.
                                                 (NITE ) NIPPON
                                                                                                                                                                                                 aggregate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 30 BP; 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 8; Col 19; 20pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-JAN-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                  2660 АСАЛАЛАЛАЛАЛАЛАЛА 2679
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                                                                                                                                                                                                                                                                                                                                                                            ACAAAAAAAAAAAAAAA 23
                                                                                                                                                                                                restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                        0.7%; Scillarity 100.0%; P. Conservative 0;
                                                                                                                                                                                                                                                              (first entry)
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                                               TELEGRAPH & TELEPHONE CORP
                                                                      93JP-00112515
                                                                                                93JP-00112515
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 2
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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No.
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                                                                                                                                                                                                           primer; cDNA;
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gene

expression -

by amplification

of mRNA followed

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2660 ACAAAAAAAAAAAAAAAA 2678

l Similarity 19; Conserv

Conservative

<u>,</u>

Mismatches

Indels

<u>.</u>

Gaps

0

100.0%;

Score 19; Pred. No.

DB 2; 5.9e+

Length 20;

.9e+03;

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RESULT 20
AAQ75569/c
ID AAQ75569 standard; DNA;
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Query Match
Best Local S
Matches 19
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Best Local (
                                                                           A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type clabelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                        Sequence
                                                                                                                                                                                                                      Disclosure; Page 5; 11pp; Japanese.
                                                                                                                                                                                                                                                    Analysis of cDNA and gene expression -
by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                   WPI; 1995-018287/03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                       01-NOV-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-AUG-1995
                                                                                                                                                                                                                                                                                                                                                             16-APR-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ75569;
                                                                                                                                                                                                                                                                                                                            (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                aggregate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19 BP; 0 A; 0 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 5; 11pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           by digestion with restriction enzymes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2660 АСААЛАЛАЛАЛАЛАЛА 2678
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19 ACAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19;
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                                                  20 BP; 0 A; 0 C; 1 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gene expression; reverse transcription; primer; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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/ 100.0%; Pr
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Pred. No.
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                                                                                                                                                                                                                                                                  amplification
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                                                                                                                                                                                                                                                                  mRNA followed
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               of of
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RESULT 21
AA075568/c
ID AAQ75568 standard; DNA; 20 BE
XX
AC AAQ75568;
XX
O4-AUG-1995 (first entry)
DT 04-AUG-1995 (first entry)
DT 04-AUG-1995 (first entry)
DT 04-AUG-1995 (first entry)
DE Reverse transcription primer
XX
Analysis; gene expression; ri
XX
Analysis; restriction enzym
XX
Synthetic.
XX
PD 01-NOV-1994.
XX
PN JP06303997-A.
XX
PN JP06303997-A.
XX
PN JP06303997-A.
XX
PN 16-APR-1993; 93JP-00112515
XX
PN 16-APR-1993; 93JP-00112515
XX
PN (NITE) NIPPON TELEGRAPH & T
XX
PN JP06303997-A.
XX
CX
Analysis of cDNA and gene e>
PT by digestion with restrictic
XX
PT by digestion with restrictic
XX
CC A method for the analysis of
CC double-stranded cDNAs w
CC the double-stranded cDNAs w
electrophoresing the digest
CC transcription primer; (b) d
RESULT 22
AAQ75567/c
ID AAQ75
XX AAQ75
XX AAQ75
XX AAQ75
XX AAQ75
XX AAQ75
XX AAA1
XX 
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Analysis of cDNA and gene expression - by digestion with restriction enzymes.
    01-NOV-1994
                                                                               JP06303997-A
                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                    aggregate;
                                                                                                                                                                                                                                                                      Analysis; gene expression; reverse transcription; primer; cDNA;
                                                                                                                                                                                                                                                                                                                                                         Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                        04-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ75567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 5; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity es 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 АСАЛАЛАЛАЛАЛАЛАЛА 2678
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20 BP; 1 A; 0 C; 1 G; 18 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      etandard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ACAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ACAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                         restriction enzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                expression; reverse triction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            93JP-00112515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.7%; Dr
100.0%; Pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 19; DB; Pred. No. 5.9
                                                                                                                                                                                                                                         88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 transcription; primer; cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               by amplification of mRNA followed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .9e+03
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RESULT 23
AAT04917/
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Best Local S
Matches 19
                    New stem cell factor polypeptide(s) - primitive progenitor cells, esp. for t
                                                                                                                                                                16-OCT-1989;
11-JUN-1990;
                                                                                                                                                                                                                                                  EP676470-A1
                                                                                                                                                                                                                                                                                               Stem cell factor; progenitor; haematopoiesis; SCF; anaemia; thrombocytopenia; leucopenia; AIDS; immunodeficiency; bone graft; transplant; neoplasia; myelosuppression; bone marrow; ss.
                                                                                                                                                                                                                                                                                                                                                                        25-MAR-2003
15-MAY-1996
                                                                                                                                                                                                                                                                                                                                                                                                            AAT04917;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in separate lanes. The electrophoresing the digested aggregate of cDNAs in separate lanes.
                                                         WPI;
                                                                                 Zsebo
                                                                                                                                         24-AUG-1990;
28-SEP-1990;
                                                                                                                                                                                                     04-OCT-1990;
                                                                                                                                                                                                                             11-OCT-1995.
                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                Mammalian stem cell factor
                                                                                                                                                                                                                                                                                                                                                                                                                                   AAT04917
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Analysis of cDNA and gene expression - by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1995-018287/03.
                                                                                                      (AMGE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2660 АСАААААААААААААА
                                                         1995-346090/45
                                                                                3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 can be used to analyse gene expression rapidly and easily
                                                                                                       AMGEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                   standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ACAAAAAAAAAAAAAAA 1
                                                                              Suggs SV,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.7%;
llarity 100.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 0 A; 0 C; 2 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 5; 11pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                        (revised)
(first en
                                                                                                       INC.
                                                                                                                           89US-00422383.
90US-00537198.
90US-00573616.
90WO-US005548.
90US-00589701.
                                                                                                                                                                                                     95EP-00105391
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             93JP-00112515
                                                                                                                                                                                                                                                                                                                                                                                                                                    CDNA;
                                                                                                                                                                                                                                                                                                                                                                        entry)
                                                                                Bosselman
                                                                                                                                                                                                                                                                                                                                                                                                                                    20
                                                                                                                                                                                                                                                                                                                                               (SCF) cDNA oligonucleotide primer 220-3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ٥,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 19;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2678
                                                                              ₽,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                    for treating disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        No.
                                                                                Martin
                                for stimulating the growth of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 2; Le
5.9e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               by amplification of mRNA followed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                    involving
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                    blood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
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Example 3;

Fig 12C;

127pp; English.

Query Match Best Local S Matches 19

Similarity

0.7%;

Score 19; ; Pred. No.

DB 2; Le U; 0 Other;

Length 20 Indels

Conservative

0,

Mismatches

0,

0

0

Sequence

20

BP; 0 A; 0 C; 2 G; 18 T; 0

AAT04915-T04922 are oligonuclectide primers and probes used for the camplification and sequencing of mammalian stem cell factor (SCF). Non-care amplification and sequencing of mammalian stem cell factor (SCF). Non-care amplification occurring SCF and C-terminally truncated polypeptides, having a common acid sequences sufficiently duplicative of naturally occurring SCF, composition cells and primordial germ stem cells. The progenitor cells, neural stem cells and primordial germ stem cells. The composition for treating leucopenia, anaemia or cells composition for treating leucopenia, anaemia or cells composition composition for treating leucopenia, anaemia or cells composition composition. They can also cells used for treating neoplasia, nerve damage, infertility, intestinal cells used for treating neoplasia, nerve damage, infertility, intestinal cells peptides for use in detection or neutralisation of SCF in serum. SCF cells peptides for use in detection or neutralisation of SCF in serum. SCF cells be useful for the treatment of AIDS and severe combined cells immunodeficiency (SCID) states alone or in combination with other factors such as IL-7. (Updated on 25-MAR-2003 to correct PF field.)

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RESULT 24
AAA13752/c
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                                                                                                                  16-OCT-1989;
11-JUN-1990;
24-AUG-1990;
28-SEP-1990;
01-OCT-1990;
04-OCT-1990;
       Example 3;
                                 Production of subject using
                                                                                                                                                                                                                                                               Stem cell factor; SCF; haematopoietic progenitor cell; blood forming; primitive progenitor cell; haematopoietic disorder; syngeneic; allogeneic; autologous bone marrow transplant; gene therapy; transfection; haematopoietic stem cell; acute blood loss; neoplasia;
                                                                                                                                                                                                                                                                                                                                                                       3752/c
AAA13752
                                                                                                                                                                                    04-OCT-1990;
                                                              WPI; 2000-259135/23.
                                                                                                                                                                                                                       EP992579-A1
                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                              Stem cell factor universal oligonucleotide 220-3.
                                                                                                  (AMGE-) AMGEN
                                                                                                                                                                                                     12-APR-2000
                                                                                                                                                                                                                                                            cancer; ss.
                                                                                                                                                                                                                                                                                                                                    27-JUL-2000
                                                                                ₹
                                                                                                                                                                                                                                                                                                                                                                                                                     19
                                                                                                                                                                                                                                                                                                                                                                                                                                CAAAAAAAAAAAAAAAA 2679
                                                                                                                                                                                                                                                                                                                                                                      standard;
     Fig 12C; 123pp; English
                                                                             Sugge
                              hematopoietic cells suitable for administration progenitor cells and expanding the cells using s
                                                                                                 INC.
                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                 89US-00422383.
90US-00537198.
90US-00573616.
90WO-US005548.
90US-00589701.
90EP-00310899.
                                                                                                                                                                                   99EP-00122861
                                                                              sv,
                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                             Bosselmann
                                                                             RΑ,
                                                                             Martin
                                                                             Ξ
                                        6
                                       Ø
                               cell
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Enhancing efficiency of transfer of polynucleotide into a target mammalian cell in vitro, involves exposing cell that expresses a cell factor receptor to stem cell factor, and introducing polynuinto cell in vitro.

and introducing polynucleotide

a stem

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AAH41331/c
ID AAH413
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          문
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     administration to a subject. The method comprises: (a) obtaining CC haematopoietic progenitor cells from a donor; and (b) expanding the cells by adding to the cells a haematopoietically effective dose of a CC polypsptide product having at least part of the primary structural CC confirmation and one or more of the biological properties of naturally occurring stem cell factor (SCF). The method is useful for stimulating CC primitive progenitor cells including early haematopoietic progenitor CC granulocyte, lymphocyte and macrophage cells. SCF results in absolute CC increases in haematopoietic cells of both myeloid and lymphoid lineages. CC cris useful for treating haematopoietic disorders. The method is useful CC for expanding early haematopoietic progenitors in syngeneic, allogeneic CC or autiologous bone marrow transplant. SCF is useful for enhancing the CC cells. SCF is also useful for combating the myelosuppressive effects of cells. SCF is also useful for combating the myelosuppressive effects of cells. SCF is also useful for combating the myelosuppressive effects of canti-HIV drugs such as AZT and for enhancing haematopoietic recovery canti-HIV drugs such as AZT and for enhancing haematopoietic recovery complasia (cancer). The present sequence represents a universal universal consideration of the present sequence represents a universal consideration in the present invention of the present sequence represents a universal consideration in the present invention of the present sequence represents a universal consideration in the present invention of the present sequence represents a universal consideration in the present invention of the present sequence represents a universal consideration in the present invention of the present sequence represents a universal consideration in the present invention of the present sequence represents a universal consideration in the present invention of the present sequence represents a universal consideration in the present sequence represents a universal consideration in the pr
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Best Local (
                                                                                                                                                                                                                                 24-AUG-1990;
01-OCT-1990;
25-NOV-1992;
21-DEC-1993;
24-MAY-1995;
                                                                                                                                                                                                                                                                                                                             16-OCT-1989;
11-JUN-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gene
                                                                                                                                                                                                                                                                                                                                                                                    31-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                      27-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                       US6207454-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Stem
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Universal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAH41331;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAH41331 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 0 A; 0 C; 2 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                 AMGE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A method has been developed of making administration to a subject. The metho
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2661 CAAAAAAAAAAAAAAAA 2679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cell factor; SCF; stem cell factor receptor; blood cell disorder; therapy; PCR primer; mutagenesis; probe; ss.
                                                                                                                                         ₹
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19
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                                                                                                                                                                               AMGEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 stem cell factor (SCF)
                                                                                                                                         Bosselman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                             INC.
                                                                                                                                                                                                              90US-00537198.
90US-00573616.
90US-00589701.
92US-00982255.
93US-00172329.
95US-00449653.
98US-00005893.
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                                                                                                                                                                                                                                                                                                                                                                                  98US-00224681
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.7%;
                                                                                                                                         ŖΑ,
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                                                                                                                                     Suggs
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RESULT 26
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Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 of transfer of a polynucleotide (I) into a target mammalian cell (II) in vitro, comprising exposing (II) that expresses a stem cell factor (SCF) receptor to a biologically active SCF, its analogue or fragment, which induces cell proliferation, and introducing (I) to (II) in vitro.

Exposure of SCF to (II) results in increased uptake of (I) into the cell. The method is useful for enhancing the efficiency of the transfer of a polynucleotide into a target mammalian cell in vitro. The method is useful in gene therapy techniques. AAH41301 to AAH41364 and AAB98351 to AAB98390 represent sequences used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              blood
The present sequence for universal PCR primer 220-3 is 1 of 8 universal oligonucleotides (AASO4110-AASO4117) used in the isolation of the human SCF (stem cell factor) cDNA sequence. The present invention relates to novel stem cell factors (AAU02453-AAU02458, AAU02460, AAU02461) and the
                                                                                                                                            Novel nucleic acids disorders involving disease, Kala azar,
                                                                                                                                                                                                                                                                                                                 (ZSEB/)
(BOSS/)
(SUGG/)
(MART/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16-OCT-1989;
11-JUN-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human
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01-OCT-1990;
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                                                                                                                                                                                                                                   2001-298941/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; stem cell factor; SCF; early haematopoietic progenitor disorder; leukaemia; Hodgkin's disease; lymphoma; splenc
                                                                                                                                                                                                                                                                            <u>$</u>
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BOSSELMAN .
SUGGS S V.
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90US-00537198.
90US-00573616.
90US-00589701.
93US-00172329.
                                                                                                                                                                                                                                                                                                                   FH.
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g blood cells, e.,
anemia and sept
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                                                                                                        English
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cDNA universal PCR
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                                                                                                                                               septicemia.
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                                                                                                                                                                                                                                                                        Martin
                                                                                                                                                                leukemia,
                                                                                                                                                                                        factor useful for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 4; 1
5.9e+03
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                                                                                                                                                                  splenomegaly,
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on disorder;
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                                                                                                                                                                    Hodgkin's
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RESULT 27
AAF89091/c
ID AAF89091;
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AC AAF89091;
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AC AAF89091;
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DE AAF89091;
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DE Mammalian stem cell:
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DE Mammalian rat; mammal;
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DE Mammalian rat; mammal;
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DE Mammalia.
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DE Mammalia.
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DE Mammalia.
XX
W Gene therapy; haemat:
XX
V Gevere combined immu
XX
PP 09-NOV-1994; 94US-
PR 11-JUN-1990; 90US-
PR 11-JUN-1990; 90US-
PR 25-NOV-1992; 92US-
PR 25-NOV-1990; 90US-
PR 25
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                                                                                                                                                                                                                                                                                          Novel isolated non-human mammalian stem cell factor polypeptide stimulating growth of early hematopoietic progenitor cells, useful fo treating aplastic anemia, lymphoma, Letterer-Siwe disease, Kala azar,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; rat; mammal; stem cell factor; SCF; cell growth stimulation; gene therapy; haematopoietic disorder; aplastic anaemia; leukaemia; neurological damage; intestinal damage; infertility; AIDS; SCID; severe combined immunodeficiency; PCR primer; ss.
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The present invention provides the protein and coding sequences of mammalian stem cell factors (SCFs). These are capable of stimulating to growth of early haematopoietic progenitor cells, neural stem cells and primordial germ stem cells. The sequences are useful in the treatment leukaemias, haematopoietic disorders, aplastic anaemia, paroxysmal nocturnal haemoglobinuria, malaria, pigmentation disorders, neurologic and intestinal damage, infertility, AIDS and severe combined

neurological

and the of.

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RESULT 28
ARH23889/c
ID AAH23889/c
AR AAH2388
XX AAH238
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XX Human;
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The present sequence for universal PCR primer 220-3 is 1 of 8 universal CC oligonucleotides (AAH23888-AAH23895) used in the isolation of the human CS CF (stem cell factor) cDNA sequence. The present invention relates to CC novel stem cell factors (AAB73561-AAB73568) AAB73571-AAB73576) and the CC including early haematopoletic progenitor cells. The invention also CC including early haematopoletic progenitor cells. The invention also CC describes SCF peptides (AAB73578-AAB73597) and the oligonucleotides CC (AAH23859-AAH23887) used in the isolation of human and rat SCF sequences. CC The polynucleotide encoding SCF is useful for producing SCF and useful in CC gene therapy. It is useful for treating disorders involving blood cells CC such as myelofibrosis, metastatic carcinoma, acute leukaemia, multiple CC congestive splenomegaly, Kala azar, sarcoidosis, military tuberculosis, CC disseminated fungus disease, lymphoma, Gaucher's disease, anaemia, CC disseminated fungus disease, pulminating septicaemia, malaria, vitamin CC and folic acid deficiency, pyridoxine deficiency, and CC hypopigmentation disorders such as piebaldism and vitiligo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-OCT-1989;
11-JUN-1990;
24-AUG-1990;
01-OCT-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 growth of early
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New stem
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-256683/26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        anemia,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-APR-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-NOV-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              blood disorder; leukaemia; Hodgkin's disease; lymphoma; splenomegaly;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; stem cell factor; SCF; early haematopoietic progenitor cell;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAH23889;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAH23889 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2661 САААААААААААААААА 2679
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SCF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     m cell factor polypeptides and their analogs which stimulate
of early hematopoietic progenitors, useful for treating aplastic
carcinoma, multiple myeloma, vitiligo, kala azar, Hodgkin's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                l Similarity
19; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bosselman
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                12C; 166pp;
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90US-00589701.
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90US-00537198.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cell factor)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cDNA universal PCR primer 220-3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 19;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Martin FH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 4;
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CC The present sequence for universal PCR primer 220-3 is 1 of 8 universal CC oligonucleotides (AASO4211-AASO4218) used in the isolation of the human CC SCF (stem cell factor) cDNA sequence. The present invention relates to CC novel stem cell factors (AAU02761-AAU02767, AAU027797) and the polynucleotides encoding them. SCF stimulate primitive progenitor CC cells including early haematopoletic progenitor cells. The invention also CC cells including early haematopoletic progenitor cells. The invention also CC (AASO4182-AASO4210) used in the isolation of human and rat SCF sequences. CC The polynucleotide encoding SCF is useful for producing SCF and useful in CC gene therapy. It is useful for treating disorders involving blood cells such as myelofibrosis, metastatic carcinoma, acute leukaemia, multiple CC unyeloma, Hodgkin's disease, lymphoma, Gaucher's disease, anaemia, CC disseminated fungus disease, Plymphoma, Gaucher's disease, anaemia, CC disseminated fungus disease, Fulminating septicemia, malaria, vitamin B12 CC and folic acid deficiency, pyridoxine deficiency, and hypopigmentation CC disorders such as piebaldism and vitiligo
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AAS04212/
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01-OCT-1990;
25-NOV-1992;
                                                                                                                                                                                                                                                                        Example 3; Fig 12C; 167pp; English
                                                                                                                                                                                                                                                                                               Isolated DNA sequence, encoding polypeptide product useful stimulating growth of early hematopoietic progenitor cells.
                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-281051/29
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11-JUN-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US6218148-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               blood disorder; leukaemia; Hodgkin's disease; lymphoma; splenomegal
anaemia; Kala azar; septicemia; malaria; hypopigmentation disorder;
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90US-00589701.
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Pred. No.
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Query Match Best Local (Matches

Similarity

0.7%; 5c. 100.0%; Pr

Score 19; DB; Pred. No. 5.9

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Length 20

Indels

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Gaps

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Conservative

Sequence 20 BP; 0 A;

0 C; 2 G; 18 T; 0 U; 0 Other;

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RESULT 30
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Query Match
Best Local S
Matches 19
                                                                           The present sequence for universal PCR primer 220-3 is 1 of 19 PCR primers (AAS10435-AAS10435) used to amplify various portions of the human SCF cDNA sequence. The sequence is described in an invention relating to novel stem cell factors, the polynucleotides encoding them and methods for producing the stem cell factors. The methods involve increasing the number of early haematopoietic progenitor cells in human peripheral blood by administering a haematopoietically effective human stem cell factor polypeptide. The methods are useful for the treatment of blood disorders, including myelofibrosis, myelosclerosis, osteopetrosis, metastatic carcinoma, acute leukaemia, multiple myeloma, Hodgkin's disease, lymphoma, Gaucher's disease, Niemann-Pick disease, refractory anaemia, malaria, vitamin B12 and folt acid deficiency, hypopigmentation disorders i.e. piebaldism and viral induced disorders, including AIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; stem cell factor; SCF; haematopoietic progenitor cell; blood disorder; Hodgkin's disease; vitamin B12; folic acid deficiency; hypopigmentation disorder; viral disorder; AIDS; PCR primer; ss.
                                                                                                                                                                                                                                                                       Example 3; Fig 12C; 210pp; English.
                                                                                                                                                                                                                                                                                                  Increasing the number of early hematopoietic progenitor cells in the peripheral blood useful for the treatment of blood disorders including Hodgkin's disease comprises the administration of human stem cell factor.
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                                                      Sequence
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(BOSS/)
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11-JUN-1990;
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25-NOV-1992;
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BOSSELMAN R
SUGGS S V.
              Similarity
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0.7%;
ilarity 100.0%;
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90US-00537198.
90US-00573616.
90US-00589701.
91US-00684535.
92US-00982255.
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Pred. No.
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RESULT 31
AAD35464/c
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                                                                                                                                                                                                                                                                                      WPI; 2002-350789/38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Letterer-Siwe disease; refractory erythroblastic anaemia; Kala azar; Di Guglielmo syndrome; congestive splenomegaly; splenic pancytopaenia; disseminated fungus disease; Fulminating septicaemia; piebaldism; AIDS acquired immune deficiency syndrome; malaria; military tuberculosis; pyridoxine deficiency; vitamin B12 deficiency; folic acid deficiency; piamond Blackfan anaemia; hypopigmentation disorder; vitiligo; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rat
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(BOSS/)
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SUGGS S V.
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                                                                                                                                                                                                                                                                                                         SV,
                                                                                                                                                                                                                                                                                                          Martin
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AIDS;

Novel non-naturally-occurring stem cell factor polypeptide, useful treating leucopenia, thrombocytopenia, anemia and for enhancing engraftment of bone marrow during transplantation in a mammal.

Example 3; Fig 12C; 217pp; English.

The present invention relates to novel non-naturally-occurring stem cell CC factor (SCF) polypeptides having an amino acid sequence sufficiently CC duplicative of that of naturally-occurring SCF to allow possession of CC haematopoietic biological activity of naturally occurring SCF. Sequences CC of the invention are useful for treating leucopaenia, thrombocytopaenia, CC anaemia and for enhancing bone marrow accovery in treatment of radiation, CC engraftment of bone marrow during transplantation in mammals and chemical CC or chemotherapeutic induced bone marrow aplasia or myelosuppression. They are also useful for treating acquired immune deficiency in a human, nerve CC damage, neoplasia, infertility, myeloproliferative disorder, intestinal CC damage in a mammal. SCF sequences are useful for preparing biologically CC active polymer polypeptide adduct, for enhancing transfection of early CC active polymer polypeptide adduct, for enhancing transfection of early CC amammal. They are useful for treating myelofibrosis, myelosclerosis, CC Hodgkin's disease, lymphoma, Gaucher's disease, netastatic carcinoma, acute leukaemia, multiple myeloma, CC Hodgkin's disease, refractory erythroblastic anaemia, Di Guglielmo CC syndrome, congestive splenomegaly, Kala azar, sarcoidosis, primary CC tuberculosis, Fulminating capticated fungus disease, malaria, military of tuberculosis, myelosclerosis, disseminated fungus disease, malaria, military vitamin Bl2 splenic pancytopaenia, disseminated fungus disease, maiaria, williary tuberculosis, Fulminating septicaemia, pyridoxine deficiency, vitamin and folic acid deficiency, Diamond Blackfan anaemia, hypopigmentation disorders such as piebaldism, AIDS (acquired immune deficiency syndrom B12

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2661 CAAAAAAAAAAAAAAAAAA 2679

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RESULT 32
ABS73848/c
ID ABS73848/c
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Best Local S
Matches 19
                              polynucleotide sequences encoding the SCFs, and methods of producing them. SCFs are involved in the blood-forming (haematopoietic) system in mammals, particularly humans. The method of the invention is useful for the production of human SCF. The stem cell factors are useful to treat disorders involving blood cells e.g. metastatic carcinoma, acute leukaemia, multiple myeloma, Hodgkin's disease, lymphoma, refractory erythroblastic anaemia, miliary tuberculosis, disseminated fungus universal oligonucleotide for SCF DNA is used in the examples of the
                                                                                                                                                                                                                                                                                                                                                                  Production of a human stem cell factor (SCF) polypeptide for disorders involving blood cells, such as leukemia, comprises mammalian cells comprising non-human SCF promoter DNA linked
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24-AUG-1990;
28-SEP-1990;
01-OCT-1990;
04-OCT-1990;
                                                                                                                                                                                                                                                                                                                Example
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11-JUN-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sugga SV,
                                                                                                                                                                                                                                                 invention relates to novel stem cell factors (SCFs)
                                                                                                                                                                                                                                                                                                                                                           human
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90US-00537198.
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90WO-US005548.
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90EP-00310899.
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Pred. No.
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to DNA
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CC haematopoietic activity and the polynucleotides encoding them. The CC haematopoietic activity and the polynucleotides encoding them. The CC polypeptides are used for treating infertility, intestinal damage, myeloproliferative disorders, leucopenia, thrombocytopenia or anaemia, cfor improving engraftment of bone marrow transplants, for enhancing bone marrow recovery after radiotherapy or chemotherapy and in treatment of carcinoma, leukaemia and military tuberculosis. The SCF polypeptides are also used to expand haematopoietic progenitor cells for transplantation and to prepare such cells for transfection with a gene. The SCF polypeptides and also as probes for mapping of the SCF gene, for identifying SCF-related diseases and as a marker for neighbouring genes. Antibodies raised against the polypeptides are useful in diagnosis and to
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ADE52460/
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Best Local
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21-DEC-1993;
24-MAY-1995;
                                                                                                                                                                        Disclosure;
                                                                                                                                                                                                        New non-natural stem
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01-OCT-1990;
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11-JUN-1990;
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                                                                                                                                                                                                                                                                                    ESSEBO K M.
BOSSELMAN SUGGS S V.
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90US-00573616.
90US-00589701.
91US-00684535.
92US-00982255.
93US-00172329.
95US-00449653.
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lso related nucleic acid
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5.9e+03
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                                                                                                                                                                                                                                                 FH
                                                                                                                                                                                          treating e.g. 1 and antibodies.
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RESULT 34
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Best Local :
                                                                                                                      initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            remove SCE
               preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of biquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition
                                                                                                                                                                                                                                                                                                                                                        Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; antisense; lung dysfunction; nasal airway dysfunction; Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory; antiallergic; antiinflammatory; antiallergic; immunosuppressive; cytostatic; gene therapy antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                            Disclosure;
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19; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sandrasagra A,
Shahabuddin
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                                                                                                                                                                                                                                                                                                            NO 3860;
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     lung allergies, or a respiratory disease or condition data for this patent is not represented in the printe
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                                                                                                                                                                                                                                        s a novel pharmaceutical composition, which has ising an oligonucleotide antisense to the region, 5' or 3' end genomic flanking regions,
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                                                         comprising oligonuclectides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonuclectides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonuclectides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic,
                                                                                                                                                                                                                                                                                                                                   Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                         WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory disease; pulmonary vasoconstriction;
          analgesic, hypotensive, immunosuppressive and cytostatic activity, beta-adrenergic agonist. The composition is useful for preventing c treating a respiratory, lung or malignant disease. The administered
                                                                                                                                                                                                                                                   This invention describes a novel composition (a) a first active agent,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              31-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection; ss; primer.
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at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nyce
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-APR-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              respiratory distress syndrome; allergic rhinitis;
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Best Local S
Matches 19
        to nucleic acids encoding a mammalian glucocorticoid receptor. The antisense oligonucleotides of the invention are useful for preventing or delaying infection, inflammation or tumour formation. The antisense oligonucleotides are also useful for treating diabetes, obesity, cardiovascular disorders, hyperlipidaemia or Cushing's syndrome. The present DNA sequence represents an antisense oligonucleotide that targets the human glucocorticoid receptor gene. NOTE: The present sequence contains 2'-methoxyethyl (2'-MOE) wings and a phosphorothicate backbone.
                                                                                                                                                                                                                                                                                                                                                  The invention
                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, obesity, cardiovascular disorder, hyperlipidemia or Cushing's syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          inflammation; tumour formation; diabetes; obesity; cardiovascular disorder; hyperlipidaemia; Cushing's syndrome; human; phosphorothioate backbone; 2'-methoxyethyl; 2'-MOE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antisense oligonucleotide; glucocorticoid receptor; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human glucocorticoid receptor-specific antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-MAY-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-MAY-2003; 2003WO-US016084.
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                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO 4182; 985pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nalseth AE;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2002US-0381857P
                                                                                                                                                                                                                                                                                      comprises an antisense oligonucleotides that are targeted ids encoding a mammalian glucocorticoid receptor. The
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Pred. No.
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. 5.9e+03;
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Best Local
                                                                                                                           The invention comprises an antisense oligonucleotides that are targeted to nucleic acids encoding a mammalian glucocorticoid receptor. The antisense oligonucleotides of the invention are useful for preventing or delaying infection, inflammation or tumour formation. The antisense oligonucleotides are also useful for treating diabetes, obesity, cardiovascular disorders, hyperlipidaemia or cushing's syndrome. The present DNA sequence represents an antisense oligonucleotide that targets the human glucocorticoid receptor gene. NOTE: The present sequence contains 2'-methoxyethyl (2'-MOE) wings and a phosphorothicate backbone.
                                                                                                     Sequence
                                                                                                                                                                                                                                                                          New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, obcardiovascular disorder, hyperlipidemia or Cushing's syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              inflammation; tumour formation; diabetes; obesity; cardiovascular disorder; hyperlipidaemia; Cushing's syndrome; human; phosphorothioate backbone; 2'-methoxyethyl; 2'-MOE.
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                        2661 САЛАЛАЛАЛАЛАЛАЛАЛА 2679
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                                                  l Similarity
19; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 glucocorticoid receptor-specific antisense oligonucleotide #4234.
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                                                                                                                       lucocorticoid receptor gene. NOTE: The present seque-methoxyethyl (2'-MOE) wings and a phosphorothioate
                                                                                                 BP; 1 A;
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5.9e+03;
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RESULT 38 ADK74647/c ID ADK74 XX

ADK74647

standard;

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RESULT 39
ADK/4442/c
ID ADK/4442/c
XX ADK/744
XX ADK/744
XX CO-MAY
DT 20-MAY
DE Chimer
XX Navl.3
KW Navl.3
KW diabet
KW infant
                                                                                                                                                                                                                                                                                                                                                                                                                          connected acid molecule encoding Navl.3, where the antisense compound connected acid molecule encoding Navl.3, where the antisense compound connected acid molecule encoding Navl.3, where the antisense compound connected with and inhibits the expression of Navl.3. The compound and composition are useful for treating a disease or condition consociated with Navl.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuroplay, chronic pain, lower back pain, confidence of the disease or condition consucrated acid from burns, migraine headache, cluster headache, mild-to-moderate confidence pain from burns, migraine headache, cluster headache, mild-to-moderate consorted but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothicate oligonucleotide with confidence represents a chimeric phosphorothicate oligonucleotide with confidence not infantile epilepsy; or ataxia. The present confidence represents a chimeric phosphorothicate oligonucleotide with confidence and a decoxy gap. Used during the antisense inhibition of thuman Navl.3 expression, the oligonucleotides are designed to target confidence and the present confidence of the human Navl.3 RNA.
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                               diabetic neuropathy;
                                               Nav1.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia;
                                                                                  Chimeric
                                                                                                                   20-MAY-2004
                                                                                                                                                                                    ADK74442 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 4; SEQ ID NO 1981; 417pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nav1.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia;
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                 infantile epilepsy;
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                                                                                phosphorothicate oligonuclectide
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Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Navl.3, where the antisense compound specifically hybridizes with and inhibits the expression of Navl.3. The compound and composition are useful for treating a disease or condition associated with Navl.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate headache; seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothicate oligonucleotide with 2,MOE wings and a deoxy gap. Used during the antisense inhibition of human Navl.3 expression, the oligonucleotides are designed to target
                                                                                                                                                                          immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
                                                                                                                                                                                                                                                                                                                                     chimeric; antisense oligonucleotide; phosphorothioate; human;
microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor;
microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic;
                                                                                                                                                    cardiovascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADM14246 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI;
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                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human mPGES-1 chimeric antisense oligonucleotide SEQ ID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                disorder;
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                                                                                                                                                    neurological disorder;
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Pred. No.
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Synthetic

Location/Qualifiers

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RESULT 41
ADM14467/c
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Best Local
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    01-JUL-2004
                                ADM14467;
                                                             ADM14467 standard;
                                                                                                                                                                                                                                                                                        antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; SEQ ID NO 433; 132pp; English.
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19; Conserv
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(first entry)
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/*tag=
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/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
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16. .20
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                                                               DNA;
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e= "2'-O-methoxyethyls"
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== "2'-O-methocyethyls"
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The present sequence represents a chimeric antisense oligonucleotide CC targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The CC human mpGES-1 gene is located no chromosome 9, more specifically to 934.3. The present invention also describes: (1) antisense compounds, CC having a sequence comprising 8-30 bp targeted to a nucleic acid encoding CC mpGES-1, which specifically hybridise with the nucleic acid encoding CC inhibits its expression; (2) a method of inhibiting the expression of CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mpGES-1. MpGES-1 chimeric CC antisense oligonuclectides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulator, and cardiovascular activities, and can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's conditionation, immunological, cardiovascular or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                              Claim 4; SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gierse
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/mod_base=
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/mote= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
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/mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                       654; 132pp; English
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Sequence 20 BP; 0 A; 0 C; 2 G; 18 T; 0 U; 0 Other;

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RESULT 42
ADP99302/c
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Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         multiple myeloma; Hodgkin's disease; lymphoma; Gaucher's disease; Niemann-Pick disease; Letterer-Siwe disease; refractory erythroblastic anaemia; Di Guglielmo syndrome; congestive splenomegaly; Kala awar; sarcoidosis; primary splenic pancytopaenia; miliary tuberculosis; diseasemiated fungus disease; Fulminating septicaemia; malaria; vitamin B12 deficiency; folic acid deficiency; pyridoxine deficiency; vitamin B12 deficiency; folic acid deficiency; pyridoxine deficiency; vitamin B12 deficiency; folic acid deficiency; pyridoxine deficiency; vitamin B12 deficiency; folic acid deficiency; piebaldism; vitaligo; neurological damage; infertility; intestinal damage; irradiation; chemotherapy; AIDS; haematopoietic recovery; acute blood loss; neoplasm; cancer; ss; PCR; primer.
The invention relates to preparing a (vertebrate) human stem cell factor (SCF) polypeptide comprising growing host cells transformed or transfected with DNA encoding a human SCF that stimulates growth of haematopoietic progenitor cells under nutrient conditions, the DNA being operatively linked to an expression control sequence, and isolating the polypeptide produced. Also included is a recombinant host cell transformed or transfected with an expression construct comprising a vertebrate SCF polypeptide-encoding DNA operatively linked to a heterologous expression regulatory sequence, permitting the expression of the vertebrate SCF polypeptide in the host cell. Disclosed as new are rat and human nucleic acids encoding SCF, SCF proteins from a number of other
                                                                                                                                                                                                                                                         Preparing a human stem cell factor (SCF) polypeptide, useful for treating hematopoietic disorders, e.g., aplastic anemia, comprises growing host cells transformed or transfected with DNA encoding a human SCF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SCF; stem cell factor; gene therapy; haematopoietic progenitor cell; aplastic anaemia; paroxysmal nocturnal haemoglobinuria; myelofibrosis; myelosclerosis; osteopetrosis; metastatic carcinoma; acute leukaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-OCT-1990;
10-APR-1991;
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11-JUN-1990;
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                                                                                                                                                                                                                             Example 3;
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21-DEC-1993;
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Conservative 0;
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90US-00589701.
91US-00684535.
92US-00982255.
93US-00172329.
95US-00449182.
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90US-00537198.
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RESULT 43
AAQ75648/c
ID AAQ756
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CC transfected prokaryotic and eukaryotic host cells, and new and useful comethods for cultured growth of such host cells capable of expression of CC SCF and its related products. The DNA sequences are also useful as CC labelled probes in isolating human genomic DNA encoding SCF, in methods CC of protein synthesis, in genetic therapy in humans and other mammals, and CC in developing transgenic mammalian species which may serve as eukaryotic consts for production of SCF and SCF products in quantity. The SCF is consequenced in developing transferic mammalian species which may serve as eukaryotic constant of screaming haematopoietic disorders, e.g., aplastic anaemia, CC useful for treating haematopoietic disorders, e.g., aplastic anaemia, CC useful for treatatic carcinoma, acute leukaemia, multiple myeloma, CC steopetrosis, metastatic carcinoma, acute leukaemia, multiple myeloma, CC syndrome, congestive splenomegaly, Kala awar, sarcoidosis, primary CC syndrome, congestive splenomegaly, Kala awar, sarcoidosis, primary CC syndrome, congestive splenomegaly, Kala awar, sarcoidosis, primary CC syndrome, congestive splenomegaly, witamia, and folic acid deficiency, CT primary conditions and vitiligo. The SCF are also useful for creating neurological damage, infertility states, intestinal damage consistion or chemotherapy, and AIDS. SCF is also useful for centuring from irradiation or chemotherapy and AIDS. SCF is also useful concerned is a universal SCF por primary used in the isolation of SCF DNA concerns a SCF DNA in the infertility of SCF DNA in t
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                                                                                                                                                                                                 WPI; 1995-018287/03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mammals and recombinantly expressed SCF protein fragments. The DNA sequences are useful for effecting the large scale synthesis of SCE variety of recombinant techniques or for generating new and useful and circular plasmid DNA vectors, new and useful transformed and
                                                                                                                                                                                                                                           (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP
                                                                                                                                                                                                                                                                                       16-APR-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           aggregate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAQ75648;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     is a universal SCF PCR primer used in the isolation of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene expression; reverse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              restriction enzyme; ss.
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                                                                                                                                                                                                                                                                                       93JP-00112515
                                                                                                                                                                                                                                                                                                                                93JP-00112515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; 0 C; 2 G; 18 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%;
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100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     transcription; primer; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 12;
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A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798)

of of

Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.

Disclosure; Page 6; 11pp; Japanese

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RESULT 44
AAQ75639/c
ID AAQ756
XX AAQ756
XX AAQ756
XX AAQ756
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XX ARST AAQ756
XX JP0630
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RESULT 45
AAQ75643/c
ID AAQ756
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Best Local S
Matches 19
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                               A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                      AAQ75643 standard;
                                                                                                                                                                                                                                                                                                                                      Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 6; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Analysis of cDNA and gene expression - by by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1995-018287/03.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   aggregate;
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                                                                                                                                                                                        2660
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19; Conserv
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                                                                                                                                                                                                                                                                                                                            BP; 0 A; 0 C; 3 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      restriction
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                    DNA;
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Pred. No.
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J. 5.9e+03;
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5.9e+03;
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                                                                                                                                                                                                                                                                              Length 21;
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RESULT 46
AAQ75646/c
ID AAQ75
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Matches
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                                                    16-APR-1993;
                                                                              01-NOV-1994.
                                                                                                         JP06303997-A
                                                                                                                                 Synthetic.
                                                                                                                                                          aggregate;
                                                                                                                                                                   Analysis; gene expression; reverse transcription; primer; cDNA;
                                                                                                                                                                                            Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                           04-AUG-1995
                                                                                                                                                                                                                                                                             646/c
AAQ75646 standard;
  (NITE ) NIPPON
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                           16-APR-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 6; llpp; Japanese.
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                                                                                                                                                         restriction enzyme;
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TELEGRAPH & TELEPHONE
                          93JP-00112515.
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                                                                                                                                                                                                                                                                             DNA;
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RESULT 47
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A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse
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RESULT 52 AAQ75640/c ID AAQ75 XX AC AAQ75

5640/c AAQ75640

standard;

DNA;

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RESULT 53
AAQ75644/c
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                                                                                                                                                                                                                                                                                                                                   Matches
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Best Local
                                                                                                                                          Analysis; ; aggregate;
                                                                                                                                                                           Reverse
                                                                                                                                                                                                                                                                                                                                                                                                     A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The electrophoresing the digested aggregate of cDNAs in seperate lanes.
WPI; 1995-018287/03
                                                                                                   JP06303997-A
                                                                                                                                                                                               04-AUG-1995
                                                                                                                                                                                                                    AAQ75644;
                                                                                                                                                                                                                                       AAQ75644 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Analysis of cDNA and by digestion with res
                                                                               01-NOV-1994
                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1995-018287/03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          aggregate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Analysis; gene expression; reverse
                                                           16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16-APR-1993;
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                  (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP
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                                                                                                                                                                         transcription
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                                                                                                                                                    gene
                                                                                                                                           gene expression; reverse transcription; primer; cDNA;
restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                              be used to analyse gene expression rapidly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            restriction
                                                                                                                                                                                                                                                                                                                                                                           BP; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Page 6; 11pp;
                                                                                                                                                                                                                                                                                                                                 0.7%; Sollarity 100.0%; I Conservative 0;
                                                                                                                                                                                             (first entry)
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                                        93JP-00112515.
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                                                                                                                                                                                                                                                                                                                                                                           0 C; 2
                                                                                                                                                                        primer used in cDNA analysis technique.
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Pred.
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Mismatches
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                                                                                                                                                                                                                                                                                                                                            DB 2; Lo
                                                                                                                                                                                                                                                                                                                                                                           U; 0 Other;
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RESULT 54
AAQ75647/c
ID AAQ75
AC AAQ75
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Best Local S
Matches 19
                                                              A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in separate lanes. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in separate lanes. The
    Sequence
                                                                                                                                                                                                                                                        Disclosure;
                                                                                                                                                                                                                                                                                                   Analysis of cDNA and gene expression by digestion with restriction enzymes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-APR-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                aggregate;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-AUG-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Analysis of a
                                                                                                                                                                                                                                                                                                                                                                                                                        (NITE ) NIPPON
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                                               be used
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                                                                                                                                                                                                                                                        Page 6; 11pp;
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    BP;
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0 C; 2 G;
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                                               analyse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      primer used in cDNA analysis technique.
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yme; ss.
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Pred. No.
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                                               expression rapidly
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                                                                                                                                                                                                                                                                                                                           amplification
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Query Match Best Local (Matches

l Similarity 19; Conserv

Conservative

0.7%; 5c-100.0%; Pr 0;

Score 19; Pred. No.

Mismatches

Indels

0

Gaps

0

5.9e+03; DB 2;

Length 21;

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RESULT 56
ADK01333/c
ID ADK013
XX
AC ADK013
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AC ADK013
XX
DT 06-MAY
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AAV35395
ID AAV35
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                                                                                                                                                                                            Best Local Similarity Matches 19; Conserva
                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                        recombinant vaccinia virus comprising of a gag gene from a retrovirus such as HIV-1 or HIV-2, fused to a DNA fragment containing an epitope region (30-300 bases in length) of a retroviral gene other than the gag gene. The gag gene may be altered so as to produce a gag protein modified from the natural sequence by the addition, deletion or substitution of at least 1 amino acid residue. The fusion gene is inserted into a region of a vaccinia virus not essential to its propagation, to give a recombinant vaccinia virus vector which is used to transform a host cell (such as Hela, Vero, VEF, rabbit kidney RK13 or human myeloma TK-143 cells). Upon culturing the host cell produces particulate structures containing the fusion gag protein. The recombinant vaccinia virus or the fusion gag protein particles may be used in the production of vaccines for protecting against infection with retroviruses such as HIV
    06-MAY-2004
                                ADK01333;
                                                            ADK01333
                                                                                                                                                                                                                                                       Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAV35388-V35414 are primers used in a method which results in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Recombinant vaccinia virus containing fusion HIB gag gene - for production in host cells of gag protein for use as vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Page 66; 84pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hypervariable region; ENV protein; vaccinia virus; gag gene; retrovirus; vaccines; infection; protection; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1998-312481/27.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (NINA-) JAPAN NAT INST INFECTIOUS DISEASES. (JAPG ) NIPPON ZEON KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9822596-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HIV-1 gag protein DNA primer #8.
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                                                          standard; DNA;
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(first entry)
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                                                            ВP
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                                                                                                                                                                                                      Score 19; DB 2; L
Pred. No. 5.9e+03;
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RESULT 57 ADK01297/c ID ADK012

ADK01297 standard; DNA; 21 BP

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Matches Query Match

Local Similarity les 19; Conser

Conservative

0.7%; 5c. / 100.0%; Prr

Score 19; DB; Pred. No. 5.9

DB 10; 5.9e+03;

Length 21; Indels

0

Gaps

0

2661 CABAABABABABABABABA 2679

Sequence 21 BP; 1 A; 0 C; 1 G; 19 T; 0 U; 0 Other;

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This invention describes a novel method for sorting single-stranded concleic acids by isolation and hybridisation of nucleic acid pools, then capture capture that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides comprise variable and non-variable regions. The capture oligonucleotides comprise variable and non-variable regions. The capture oligonucleotides comprises all possible combinations of up to 10 nucleotides to allow that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LNA) and the anchor region capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, conducting properties and especially in the form of a chip. Its surface is physical, stimulated by an electrical field or through a molecular sieve. The method is used (1) for analysis of patterns, especially in mucosal, activity of pharmaceuticals and/or nutritional compounds, e.g. food acids (anino, carboxylic or fatty acid) or their derivatives, salts and captures. The method provides rapid, inexpensive and reproducible acid sols based on comparison of fatty acid) or their derivatives, salts and capture notes; small differences in the nucleic acid pools. Since the method is based on comparison of mucleic acid pools, not individual capturing enems.
                                    capture probes used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
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cell; food additive; food supplement.
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                                 method of the invention.
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ADK01297;

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This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then creading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid metrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at cleast once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particulary locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, by stranded nucleic acids to the surface is (quasi)covalent, supramolecular, by stranded nucleic acids to the surface is (quasi)covalent, supramolecular, by stranded nucleic acids to allow a electrical filed or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair you blood, nerve or germ cells and (ii) for determining the activity of pharmaccuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and cardives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and cardives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and cardives or supplements, especially minerals, trace elements, organic acids elements, o
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germ cell; food additive; food supplement.
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19

2661 CARARARARARARARARA 2679

Query Match Best Local Matches

Similarity

100.0%; 0.7%;

Score 19; Pred. No. Mismatches

DB 10; 5.9e+03;

Length 21 Indels

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Gaps

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Sequence

22

BP; 0

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1 C; 1

G; 19

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Other

Matches

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l Similarity 19; Conserv

Conservative

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Mismatches

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Length 21; Indels

0.7%; Score 19; 100.0%; Pred. No.

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Sequence

21

BP; 2

A; 0 C; 1 G; 18 T; 0 U; 0 Other;

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RESULT 58

ADKO1335/C...

ID 01ADKO1335;

XX ADKO1335;

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XX BAY-20

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                                                      This invention describes a novel method for sorting single-stranded CC nucleic acids by isolation and hybridisation of nucleic acid pools, then CC reading out, where the nucleic acids are selectively bound using capture CC agents that are (a) immobilised on the surface of a solid matrix and (b) CC comprise variable and non-variable regions. The capture oligonuclectides CC have a 5'-invariable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region CC that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LMA) and the anchor region CC comprises a sequence of 10-50, particularly 15-25, T residues. The capture CC agents are particularly locked nucleic acids (LMA) and the anchor region CC combined to the surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, CC metal, resin, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is stranded nucleic acids to the surface is (quasi)covalent, supramolecular, CC physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, having the cacids (amino, carboxylic or fatty acid) or their derivatives, salts and conjunctives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and conjunctives or supplements, especially minerals, trace elements, organic mixtures. The method provides rapid, insxpensive and reproducible can detect very small differences in pools of nucleic acid from cells. It compounds is based on comparison of nucleic acid pools, not individual compounds is based on comparison of nucleic acid pools, not individual compounds is acid. The method of the invention.
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blood; nerve; germ cell; food additive; food supplement.
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RESULT 59 ADK01302/c ID ADK013 밁 á 28-FEB-2002; 2002DE-01008794. 28-FEB-2002; 2002DE-01008794. 04-SEP-2003 DE10208794-A1 Rattus sp. Rat DNA microarray capture oligonucleotide #22 ADK01302; blood; nerve; ss; hybridisation; 06-MAY-2004 (first entry) ADK01302 standard; DNA; 21 2661 CAAAAAAAAAAAAAAAAAA 2679 CAAAAAAAAAAAAAAA 2 tion; capture oligonucleotide; pattern; mucosal; hair root; germ cell; food additive; food supplement. ВP

(DEGS) DEGUSSA BIOACTIVES GMBH Dieck HT, Норре н;

WPI; 2003-714082/68

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable patterns and screenin and constant regions.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded CC nucleic acids by isolation and hybridisation of nucleic acid pools, then CC agents that are (a) immobilised on the surface of a solid matrix and (b) CC comprise variable and non-variable regions. The capture oligonucleotides CC least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow that comprises all possible combinations of up to 10 nucleotides to allow CC binding of particular sorts of single stranded nucleic acids. The capture capture oligonucleotides to allow comprises a sequence of 10-50, particularly 15-25. T residues. The capture capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, conducting properties and especially in the form of a chip. Its surface capture of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic cativity of pharmaceuticals and/or nutritional compounds, e.g. food acids (amino, carboxylic or fatty acid) or their derivatives, salts and captures in acids from cells. It can detect very small differences in pools of nucleic acid pools, since the method is based on comparison of nucleic acid pools, not individual can be should capture probes used in the method of the invention.

Sequence 21 BP; 0 A; 0 C; 3 G; 18 T; 0 U; 0 Other;

RESULT 60 ADK01334/c ID ADK013 밁 Matches 19; Query Match WPI; 2003-714082/68 28-FEB-2002; 2002DE-01008794. 28-FEB-2002; 2002DE-01008794. ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement. Rat DNA microarray capture oligonucleotide #54. DE10208794-A1 Rattus sp. 06-MAY-2004 (first entry) ADK01334; (DEGS) DEGUSSA BIOACTIVES GMBH. ADK01334 standard; DNA; 21 Local 2661 CAAAAAAAAAAAAAAAAA 2679 19 Similarity CAAAAAAAAAAAAAAAA 1 Conservative Dieck HT, 100.0%; Hoppe H; ВP Score 19; Pred. No. Mismatches DB 10; 5.9e+03; Length 21; Indels 0 Gaps 0

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded councleic acids by isolation and hybridisation of nucleic acid pools, then concleic acids by isolation and hybridisation of nucleic acid pools, then capture acid pools, then capture that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular steve. The method is used (i) for analysis of patterns, especially in mucosal, aciditives or supplements, especially minerals, trace elements, organic aciditives or supplements, especially minerals, trace elements, organic aciditives, insign of the complete said) or their derivatives, salts and creproducible single fine the complete pattern of all nucleic acids from cells. It can detect very small differences in pools of nucleic acid pool. Since the method is based on comparison of nucleic acid pool, since the method is based on comparison of nucleic acid pool, not individual genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent

capture probes used in

the

method

of the invention

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RESULT 61
ADX01303/c
ID ADX01303/c
ADX01303/
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CC This invention describes a novel method for sorting single-stranded CC nucleic acids by isolation and hybridisation of nucleic acid gools, then CC reading out, where the nucleic acids are selectively bound using capture CC agents that are (a) immobilised on the surface of a solid matrix and (b) CC comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region CC binding of particularly locked nucleic acids under anchor region that comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, CC metal, resin, gel, crystalline material and/or membrane, having semi-conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, having cathovylic or fatty acid) or their derivatives, salts and CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and CC mixtures. The method provides rapid, inexpensive and reproducible of continues. The method provides rapid, inexpensive and reproducible. It
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example; Page 5; 8pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sorting single-stranded nucleic acid, patterns and screening active agents,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Boekenkamp D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-FEB-2002; 2002DE-01008794.
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19; Conser
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             useful for analyzing expression uses capture agent with variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 10;
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ADK01307/c
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       connucleic acids by isolation and hybridisation of nucleic acid pools, then corrections out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at clear once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow that comprises all possible combinations of up to 10 nucleotides to allow combining of particular sorts of single stranded nucleic acids. The capture agents are particularly 10-00, particularly 15-25, Tresidues. The capture comprises a sequence of 10-50, particularly 15-25, Tresidues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, certain, resin, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, physical, strimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, activity of nbarracenticals and/or nutritional formounds as a food
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               allows imaging of the complete pattern of all nucleic acid in a cell, a can detect very small differences in the nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent
                                                                                                                                                                                                                                                                                                                                                                                                                 Example; Page 5; 8pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sorting single-stranded nucleic acid, useful for analyzing expression parterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI;
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                                                                                                                                                                                                                                                                                                                                                                            This invention describes a novel method for sorting single-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Boekenkamp D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-FEB-2002; 2002DE-01008794
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DEGUSSA BIOACTIVES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              probes used in the method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CAAAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   tion; capture oligonucleotide; pattern; mucosal;
germ cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dieck HT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                   German.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 10; I
5.9e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  <u>,</u>
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e.g.

additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible representation of differences in pools of nucleic acids from cells. It allows imaging of the complete pattern of all nucleic acid in a cell, a can detect very small differences in the nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent capture probes used in the method of the invention. and

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21 BP; 0 A; 2 C; 1 G; 18 T; 0 U; 0 Other;

Query Match Best Local 19; Similarity Conservative 0.7%; 5c. 100.0%; Pr Score 19; Pred. No. Mismatches DB 10; 5.9e+03 Length 21 Indels ٥, Gaps

0

밁 19 САЛАЛАЛАЛАЛАЛАЛАЛ 2679 CAAAAAAAAAAAAAAAA 1

RESULT 63 ADK01304/c ADK01304; ADK01304 standard; DNA; 21

06-MAY-2004

(first

entry)

Rat DNA microarray capture oligonucleotide #24.

hybridisation; od; nerve; germ capture oligonucleotide; pattern; mucosal; hair root; cell; food additive; food supplement. supplement

Rattus

DE10208794-A1

04-SEP-2003.

28-FEB-2002; 2002DE-01008794

28-FEB-2002; 2002DE-01008794.

(DEGS) DEGUSSA BIOACTIVES GMBH.

Boekenkamp D, Dieck HT, Hoppe H;

WPI; 2003-714082/68

patterns constant single-stranded nucleic acid, useful for analyzing expression s and screening active agents, uses capture agent with variable

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded CC nucleic acids by isolation and hybridisation of nucleic acid pools, then compared the problem of a selectively bound using capture in mobilised on the surface of a solid matrix and (b) CC comprise variable and non-variable regions. The capture oligonuclectides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (LNA) and the anchor region CC comprises a sequence of 10-50, particularly 15-25, T residues. The CC capture oligonucleotides are bictinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single surface is remarked murlair acids to the surface is particularly a layer of (bio)molecular filaments and binding of single to the surface is (quasi)covalent,

> Query Match Best Local S Matches 19 physical, stimulated by an electrical field or through a molecular sieve The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible representation of differences in pools of nucleic acids from cells. It allows imaging of the complete pattern of all nucleic acid in a cell, an method is based on comparison of nucleic acid pools. Since the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent capture probes used in the method of the invention. Sequence 21 Similarity BP; 0 Conservative A; 0 100.0%; 0.7%; C; 2 0 Ģ. Score 19; Pred. No. 19 T; 0 U; 0 Other; Mismatches DB 10; 5.9e+03; Length 21 a molecular sieve. <u>.</u> Gaps and 0

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RESULT 64

ADK01306 standard; DNA; 21 ₽P

ADK01306;

06-MAY-2004 (first entry)

Rat DNA microarray capture oligonucleotide #26.

ss; hybridisation; blood; nerve; germ germ capture oligonucleotide; pattern; mucosal; hair root; cell; food additive; food supplement.

Rattus gp.

DE10208794-A1

04-SEP-2003

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

(DEGS) DEGUSSA BIOACTIVES GMBH

WPI; 2003-714082/68

Boekenkamp D,

Dieck HT, Hoppe H;

patterns and const Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable

Example; Page 5; 8pp; German.

ADKOl306/c
ID ADKOl3
XX
AC ADKOl3
XX
AC ADKOl3
XX
DT 06-MAY
XX
DE Rat DN
XX
DE Rat DN
XX
COS Rattue
XX
OS Rattue
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DE102(
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DE102(
XX
PF 28-FEI
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PF 28-FEI
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PF 28-FEI
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PF 28-FEI
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PF Boeke:
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PF Boeke:
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DEGS
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This
CC This
CC This
CC That
CC Compri
CC Capent
CC CAPEN
CC CA This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass,

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RESULT 65
ADK01299/c
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cc conducting properties and especially in the form of a chip. Its surface
cs is particularly a layer of (bio)molecular filaments and binding of single
cc stranded nucleic acids to the surface is (quasi)covalent, supramolecular,
cc physical, stimulated by an electrical field or through a molecular sieve.
cc The method is used (i) for analysis of patterns, especially in mucosal,
cc hair root, blood, nerve or germ cells and (ii) for determining the
cc activity of pharmaceuticals and/or nutritional compounds, e.g. food
cc additives or supplements, especially minerals, trace elements, organic
cc acids (amino, carboxylic or fatty acid) or their derivatives, salts and
cc mixtures. The method provides rapid, inexpensive and reproducible
cc representation of differences in pools of nucleic acids from cells. It
cc allows imaging of the complete pattern of all nucleic acid in a cell, and
cc can detect very small differences in the nucleic acid mool. Since the
cc method is based on comparison of nucleic acid pools, not individual
cc capture probes used in the method of the invention.
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Best Local S
Matches 19
This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture
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Sorting s

single-stranded nucleic acid,
s and screening active agents,

useful for analyzing expression uses capture agent with variable

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constant

regions.

Example;

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gpp;

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This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b)

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CC agents are particularly locked nucleic acids (LNA) and the anchor region CC comprises a sequence of 10-50, particularly 15-25. T residues. The CC capture oligonucleotides are biotinnylated and immobilised on a surface by CC interaction with streptavidin. The matrix is of plastic, ceramic, glass, CC metal, resin, gel, crystalline material and/or membrane, having semi-CC conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single CC stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. CC The method is used (i) for analysis of patterns, especially in mucosal, CC hair root, blood, nerve or germ cells and (ii) for determining the CC activity of pharmaceuticals and/or nutritional compounds, e.g. food CC additives or supplements, especially minerals, trace elements, organic CC mixtures. The method provides rapid, inexpensive and reproducible CC representation of differences in pools of nucleic acids from cells. It CC allows inaging of the complete pattern of all nucleic acid made cell, and CC anish based on comparison of nucleic acid pool. Since the complete pattern of all nucleic acid pool. Since the matrix miniaturisation is possible. ADX01281-ADX01344 represent comparison of the complete pattern of the nucleic acid pool.
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RESULT 67
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                                                                       Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.
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                                                                                                                                                                                           Boekenkamp D,
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This invention describes a novel method for sorting single-stranded

Example; Page 5; 8pp; German.

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ADK01300/c
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Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.
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                                                                                                                                            Dieck
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Pred. No. 5.9e+03;
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RESULT 69
ADK01305/c
ID ADK013
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CC that comprises all possible combinations of up to 10 nucleotides to allow the agents are particular sorts of single stranded nucleic acids. The capture CC agents are particularly locked nucleic acids (LNA) and the anchor region CC comprises a sequence of 10-50, particularly 15-25, T residues. The CC capture oligonucleotides are biotinylated and immobilised on a surface by CC interaction with streptavidin. The matrix is of plastic, ceramic, glass, CC metal, resin, gel, crystalline material and/or membrane, having semi-cc conducting properties and especially in the form of a chip. Its surface stranded nucleic acids to the surface is (quasi)covalent, supramolecular, CC physical, stimulated by an electrical field or through a molecular sieve. CC The method is used (i) for analysis of patterns, especially in mucosal, CC matrix of pharmaceuticals and/or nutritional compounds, e.g. food CC additives or supplements, especially minerals, trace elements, organic caids (amino, carboxylic or fatty acid) or their derivatives, salts and CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and callows imaging of the complete spid, inexpensive and reproducible callows imaging of the complete pattern of all nucleic acid from cells. It CC allows imaging on comparison of nucleic acid pool. Since the CC andetect very small differences in the nucleic acid pool. Since the CC andetect very small differences in the nucleic acid pool. Since the CC andetect very small and the marked of the invention.
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Best Local Similarity
WPI; 2003-714082/68
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADK01305 standard; DNA; 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region.
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                                       Boekenkamp D,
                                                                                                                                                                                                                                                                      DE10208794-A1
                                                                                                                                                                                                                                                                                                                                                                               ss; hybridisation;
                                                                                                                                                                                                                                                                                                                                                                                                                           Rat DNA microarray capture oligonucleotide
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                                                                                                                               28-FEB-2002; 2002DE-01008794
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                                                                                      DEGUSSA BIOACTIVES GMBH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CAAAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                        tion; capture oligonucleotide; pattern; mucosal;
germ cell; food additive; food supplement.
                                       Dieck HT,
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Pred. No.
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. 5.9e+03
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XFFFX#X95050505050505050505050505050505050
                                                                            Comprise variable and non-variable regions. The capture oligonuclectides to comprise variable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow cCC binding of particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (LNA) and the anchor region CC comprises a sequence of 10-50, particularly 15-25, T residues. The CC capture oligonucleotides are biotinylated and immobilised on a surface by conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, CC physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, cardistives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and carboxylic or fatty acid) or their derivatives, salts and carboxylic or fatty acid) or their derivatives, salts and carboxylic or fatty acid) or their derivatives, the coll, and cell, and cell, and cell, and cell, and cell acids of the comparison of nucleic acid pool. Since the cells and only on the method of the method of the invention
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Query Match
Best Local Similarity
Matches 19; Conser 2661 CAAAAAAAAAAAAAAAAA 2679 0.7%; ilarity 100.0%; Conservative (0 Score 19; Pred. No. Mismatches DB 10; 1 5.9e+03; 0 Length 21; Indels 0 Gaps 0

Sequence 21

capture

probes used in the method of the invention.

BP; 1 A; 1 C; 1 G; 18 T; 0 U; 0 Other;

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          RESULT 70
ADK01342/c
ADK01342 standard; DNA; 21
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                                                CAAAAAAAAAAAAAAAAA 1
 ВP
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06-MAY-2004

(first entry)

28-FEB-2002; 2002DE-01008794. 04-SEP-2003 DE10208794-A1 ss; hybridisation; capture oligonucleotide; pattern; mucosal; blood; nerve; germ cell; food additive; food supplement. Rat DNA microarray capture oligonucleotide #62. 28-FEB-2002; 2002DE-01008794 Rattus ďB

(DEGS) DEGUSSA BIOACTIVES GMBH

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CC This invention describes a novel method for sorting single-stranded CC nucleic acids by isolation and hybridisation of nucleic acid pools, then creating out, where the nucleic acids are selectively bound using capture CC agents that are (a) immobilised on the surface of a solid matrix and (b) CC comprise variable and non-variable regions. The capture oligonucleotides CC have a 5'-invariable and non-variable region, the complement of which is present at cleast once in each nucleic acid and a 3'-variable, discriminatory region CC that comprises all possible combinations of up to 10 nucleotides to allow Sinding of particularly locked nucleic acids (LNA) and the anchor region CC comprises as sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by CC capture oligonucleotides are biotinylated and immobilised on a surface by CC capture oligonucleotides are biotinylated and immobilised on a surface by CC capture oligonucleotides are biotinylated and immobilised on a surface by CC capture oligonucleotides and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single CC physical, stimulated by an electrical field or through a molecular; of the method is used (i) for analysis of patterns, especially in mucosal, acids (maino, carboxylic or fatty acid) or their derivatives, salts and representation of differences in pools of nucleic acid from cells. It can detect very small differences in pools of nucleic acid from cells. It can detect very small differences in the nucleic acid pool. Since the genes, matrix miniaturisation is possible. AbK01281-AbK01344 represent xx
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Best Local
                                                                                                     Rattus
                                                                                                                                                            ss; hybridisation,
                                                                                                                                                                                     Rat DNA microarray capture oligonucleotide #28
                                                                                                                                                                                                                                06-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                        ADK01308 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example; Page 6; 8pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
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                                                                                                                                      germ
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                                                                                                                                  capture oligonucleotide; pattern; mucosal; hair root;
cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.7%; Score 19;
100.0%; Pred. No.
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This invention describes a novel method for sorting single-stranded CC nucleic acids by isolation and hybridisation of nucleic acid pools, then computed acids out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) CC comprise variable and non-variable regions. The capture oligonucleotides CC have a 5'-invariable and non-variable region. The complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region CC that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bo) molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, the method is used (i) for analysis of patterns, especially in mucosal, cardistives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and cardis (amino, carboxylic or fatty acid) or their derivatives, salts and cardis (amino, carboxylic or fatty acid) or their derivatives, salts and cardis (amino, carboxylic or fatty acid) or their derivatives, salts and cardis (amino, carboxylic or fatty acid) or their derivatives, salts and cardis (amino, carboxylic or fatty acid) or their derivatives, salts and cardis (amino, carboxylic or fatty acid) or their derivatives, salts and cardis (amino, carboxylic or fatty acid) or their derivatives, salts and cardis or mucleic acid pool. Since the capture probes used in the method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example; Page 5; 8pp; German.
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Sequence 21 BP; 0 A; 1 C; 1 G; 19 T; 0 U; 0 Other;

Matches Query Match Best Local Similarity 19; Conservative 100.0%; 0.7%; ٥, Score 19; ; Pred. No. Mismatches DB 10; 5.9e+03; Length 21; Indels <u>,</u> Gaps

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닭 S 2661 CAAAAAAAAAAAAAAAAAA 2679 19 CAAAAAAAAAAAAAAAA 1

ABA93238 RESULT 72 PolyA adaptor oligonucleotide SEQ ID NO:1. Synthetic Detection; 18-APR-2002 ABA93238; ABA93238 standard; DNA; 22 comparative detection; adaptor; (first entry)

JP2001333800-A

28-FEB-2002; 2002DE-01008794

04-SEP-2003 DE10208794-A1

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Best Local
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03-AUG-1995
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                                                        13-MAY-1993;
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                                                                                                                                                                                                                                                                                      LCR oligo
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                (USSH ) US DEPT
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                                                                                                                                                                                                                                                                                                                                                                                                                        standard;
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llarity 100.0%;
Conservative (
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                                                      93US-00061694.
                                                                                            94WO-US005407.
              HEALTH & HUMAN SERVICES
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62942 are examples of synthetic immunoreactive peptides. a method for detecting an antigen in a subject. The binding the antigen to a solid support and then reacting

(L) bound to an

support and an oligo; rer

removing any

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immunoreactive ligand

AAR62941 and AAR62942

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Example; Page 13; 34pp; English

diagnosing hepatitis C

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infection.

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                                                                                          Solid phase immunoassay using oligo:nucleotide as label -conjugates of oligo:nucleotide coupled to antigenic peption
                                                                                                                               WPI; 1995-006819/01.
                                                                                                                                                     Fields HA, Khudyakov
                                                                                                                                                                                                     13-MAY-1993;
                                                                                                                                                                                                                                                    24-NOV-1994.
                                                                                                                                                                                                                                                                            WO9426932-A1
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03-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Solid phase immunoassay using oligo:nucleotide as laconjugates of oligo:nucleotide coupled to antigenic diagnosing hepatitis C or E virus infection.
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                                                                                                                                                                                                                                                                                                                            Synthetic oligo;
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RESULT 75 AAZ00877/c Ş 밁 SSSSSSSSSSS Query Match Best Local S Matches 19 PG1 gene; biallelic marker; PCR primer; PG1-related biallelic marker; cancer; prostate cancer; diagnosis; therapy; prostate specific antigen; PSA; human; ss. unreacted L, and then detecting the presence of the oligo. A similar method can be used to detect Abs, in which case the ligand is an oligo-labelled Ag. The use of an amplifiable oligo as the label allows Ag or Ab to be detected at very low levels. An examplary oligi is AAQ75024 which can be covalently attached by the 5'- terminus to the N- or C-terminal of a synthetic peptide. For LCR using oligo AAZ75024, oligos 1-4 (see AAQ75027-Q75030) can be used. (Updated on 25-MAR-2003 to correct PN PCR primer PGRT32 27-SEP-1999 AAZ00877 standard; DNA; 24 AAZ00877; Sequence 23 BP; 0 A; 2661 CAAAAAAAAAAAAAAAAAA 2679 19 САЛАЛАЛАЛАЛАЛАЛАЛА 1 19; Similarity 100.0%, p (first entry) for PG1 coding sequence. 0 C; 4 G; 1 T; 18 U; 0 Other; 0.7%; ₽P Score 19; Mismatches ; DB 2; Lo 0; Length 23; Indels 0, Gaps 0,

Homo Synthetic. sapiens.

WO9932644-A2

01-JUL-1999

22-DEC-1998; 98WO-IB002133

22-DEC-1997; 09-SEP-1998; 97US-00996306. 98US-0099658P.

(GEST) GENSET.

Cohen D, Blumenfeld M, Chumakov I, Bougueleret L;

WPI; 1999-405178/34.

Use of a prostate cancer associated gene and biallelic markers derived from it.

Example 6; Page 42; 385pp; English.

The invention relates to a mammalian PG1 gene and protein, and a set of CC PG1 biallelic markers. The PG1 polynucleotide and biallelic markers are CC used in a hybridisation assay, a sequencing assay, or in an allele-cC specific amplification assay for determining the identity of a nucleotide CC at a PG1-related biallelic marker. The methods can be used to detect and CC diagnosis of prostate cancer relies on prostate specific antigen (PSA) CC diagnosis of prostate cancer relies on prostate specific antigen (PSA) CC disage. However, the effectiveness of this is limited due to its CC of the organ. A need exists for both a reliable diagnostic procedure CC which would enable early-stage diagnosis, and for preventative and CC curative treatments of the disease. The PG1 gene can be used for CC detection of prostate cancer, and the risk of developing it in the CC future, and can also be used to determine therapies for the disease

Sequence 24 BP; 3 A; 0 C; 1 <u>ი</u> 20 H 0 U; 0 Other

> Matches Query Match 0.7%; Best Local Similarity 100.0%; Conservative ٥. Score 19; Pred. No. Mismatches 5. 5. .8e+03; Length 24;

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2661 CAAAAAAAAAAAAAAAAAA 2679

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Search completed: February Job time : 1262.01 secs 2, 2005, 16:08:58

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1 21 0.8 25 1 ALO48782 2 21 0.8 29 1 ALO39119 3 20 0.7 21 8 MS558161 4 20 0.7 21 8 MZ528161 5 20 0.7 23 1 ALO38592 7 20 0.7 23 1 ALO38699 8 20 0.7 23 1 ALO38688 9 20 0.7 23 1 ALO38688 9 20 0.7 23 1 ALO38699 10 20 0.7 23 5 MX559898 10 20 0.7 28 6 CF336199 11 20 0.7 28 7 T52836 12 20 0.7 28 8 AZ654007 13 20 0.7 30 1 ALO38731 14 20 0.7 30 1 ALO38731 15 20 0.7 30 5 MX553986 16 20 0.7 30 6 CF298396 17 20 0.7 30 8 AZ458127 19 0.7 20 1 ALO38460 20 19 0.7 20 1 ALO38460 21 19 0.7 20 1 ALO3845727 23 19 0.7 20 1 ALO3845727 24 19 0.7 20 1 ALO38470727 25 19 0.7 20 1 ALO38470727 26 CF29820002	score greater than or equal to the score of the and is derived by analysis of the total score d and is derived by analysis of the total score d SUMMARIES Result Query No. Score Match Length DB ID	EST:* 1: gb est1:* 2: gb est2:* 3: gb_htc:* 4: gb est3:* 5: gb est4:* 6: gb est4:* 6: gb est4:* 9: gb est5:* 9: gb est5:*	um DB seq length: 0 um DB seq length: 30	Word size : 0 Total number of hits satisfying chosen parameters:	Scoring table: OLIGO_NUC Gapop 60.0 , Gapext 60.0 Searched: 32822875 seqs, 18219865908 residues	Title: US-10-048-046-1 Perfect score: 2679 Sequence: 1 aagaattcggcacgaggccgacaaa	on: February 2, 2005, 15:07:30 ; Searc (without 11866.38	GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen OM nucleic - nucleic search, using sw model
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AUTHORS
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AL048782
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KEYWORDS
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                                                                                                                        ORGANISM
            Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
1 (bases 1 to 25)
Koehrer, K., Beyer, A., Mewes, H.
EST (Koehrer, et al.)
                                                                                                                                                                                                               AL048782
DKFZp5660013 r1 566 (synonym:
DKFZp5660013, mRNA sequence.
                                                                                                                      Homo sapiens
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: hfkd2) Homo sapiens cDNA clone
                                                                           Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                  Gassenhuber, J. and Wiemann, S.
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DKFZp566J034, mRNA sequence.
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Unpublished (1999)
Contact: MIPS
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                                                                                                                                                                                                                                  0.8%; Solilarity 100.0%; I
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/db_xref="taxon:9606"
/clone="DKFZp5660013"
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 29)
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    Location/Qualifiers
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Location/Qualifiers
                                                                                                                                                                                                                                                                            /organism="Homo sapiens"
/mol type="mRNA"
/mb ref="mRNA"
/db ref="taxon:9606"
/clome="DKFZp566J034"
/tissue_type=""kidney"
/dev stage="fetal"
/lab host="x1-zblue"
/clome_lib="566 (synonym: hfkd2)"
/note="Vector: pAMP1; Site_1: NotI; Site_2: Sa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /tissue_type="kidney"
/dev_stage="fetal"
/dev_stage="fetal"
/lab_host="X1-2blue"
/clone_lib="566 (synonym: hfkd2)"
/note="Vector: pAMP1; Site_1: NotI; Site_2: SalI"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mewes, H.W., Gassenhuber, J. and
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gut
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RESULT 4
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Matches 20; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Prof. M.J.Lehane Coll. School of Biological Sciences, University of Wales, Bargor LL57 2UW All clones with suffix old are
                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 21)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          All clones with suffix qlc are reverse primer reads starting at 5' end of the cDNA all plc reads are from the 3' end.
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Pathogen Sequencing Unit
The Sanger Institute The Wellcome Trust Genome Campus
Hinxton, Cambridge, CB10 1SA, UK
Request for clones, please contact: Mike Lehane
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Glossina morsitans morsitans
Glossina morsitans
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22881942
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1 (bases 1 to 21)
                                                                                                                                                                                                  Mus musculus
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               plasmid inserts
Unpublished (2000)
                                                 Mouse whole genome scaffolding with paired end
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone="Tse37a05_plc"
/tissue_type="adult infected gu
/clone_lib="Glossina morsitans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mol_type="mRNA"
/sub_species="morsitans"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         organism="Glossina morsitans morsitans"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         note="country: Zimbabwe; EST
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100.0%; Pred. No.
B. Weiss
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S. and Hall,N.
setse fly Glossina
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                                                                                                                                                                                                                                                                                                      DKFZp566N082 rl 566 (synonym: DKFZp566N082, mRNA sequence
                                                                                                                                                   Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates; 1 (bases 1 to 23)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0045 row: M column: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                84112, USA
Tel: 801 585 5606
Fax: 801 585 7177
Ingolstaedter Landstr.1, D-8
    Location/Qualifiers
                                                         Contact:
                                                                         Unpublished (1999)
                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                             AL038397
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Seq primer: CGTTGTAAAACGACGGCCAGT
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                                                                                                                Wiemann,S.
                                                                                                                                  Ottenwaelder,B.,
                                                                                                                                                                                                                                                                           AL038397.1
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                                                                                                                                                                                                                                    Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWB42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent B. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                         MIPS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (http://www.jax.org/resources/documents/dnares/). The DN/was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"/Clone_lib="Mouse 10kb plasmid UUGCIM library"/note="Vector: pW0421v; Purified genomic DNA from M.musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Biomedical
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/strain="C57BL/6J"
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                D-85764 Neuherberg, Germany
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i: hfkd2)
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                                                                                                                                  Mewes,H.W.,
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5.6e+02;
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RESULT 7
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Best Local &
 ORGANISM
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Best Local Similarity
Matches 20; Conserv
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 Homo
                                               DKFZp566G0946_r1
DKFZp566G0946, mR
AL038609
                                  AL038609.1 GI:49682169
                                                                                    AL038609
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Mammalia; Eutheria; Primates;
1 (bases 1 to 23)
Ottenwaelder,B., Obermaier,B.,
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Location/Qualifiers
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Unpublished (1999)
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DKFZp566G1446_x1_566 (вупопут: hfkd2) Homo
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sapiens
sapiens
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ilarity 100.0%;
Conservative (
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/lab host="X1-2blue"
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/clone="DKF2p56N082"
/tlssue_type="Midney"
/dev_stage="fetal"
/lab host="xl-zblue"
/lab host="xl-zblue"
/clone_lib="566 (synonym: hfkd2)"
/note="Vector: paMP1; Site_1: NotI; Site_2:
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/mol_type="mRNA"
/db_xref="taxon:9606"
           (human)
                                                                                                                                                                                                         0.7%; Score 20;
100.0%; Pred. No.
                                                                     566 (synonym:
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                                                         sequence.
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                                                                     hfkd2) Homo
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J. 5.6e+02;
O;
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5.6e+02;
hes 0;
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CDNA clone
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Best Local :
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                                                                 Similarity
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Ottenwaelder,B., O
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2660 АСААЛАЛАЛАЛАЛАЛАЛ 2679
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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    Location/Qualifiers
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1 (bases 1 to 23)
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Location/Qualifiers
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                                                                 0.7%; Score 20;
larity 100.0%; Pred. No.
Conservative 0; Mismatch
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/dev stage="fetal"
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                                                                                                                                                       /organism="Homo sapiens"
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/dev_stage="fetal"
/lab_host="X1-2blue"
/clone_lib="566 (synonym: hfkd2)"
/note="Wector: pAMP1; Site_1: NotI; Si
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5.6e+02;
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CF336199/c
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BX559898/c
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Pathogen Sequencing Unit
The Sanger Institute The Wellcome Trust Genome Campus
Hinxton, Cambridge, CB10 1SA, UK
Request for clones, please contact: Mike Lehane
Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Cyza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.

1 (bases 1 to 26)
                                                                                                                                                                           JMT--06-C18.g1 AtJMT-overexpressing transgenic rice library (JMT) Oryza sativa (japonica cultivar-group) JMT--06-C18, mRNA sequence.
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Bangor LL57 2UW
All clones with suff
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Genome Biol. 4 (10), R63 (2003)
22881942
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EX559898 23 bp mRNA linear EST 10-
EX559898 Glossina morsitans morsitans adult infected gut G
morsitans morsitans cDNA clone Tse46d05_plc, mRNA sequence
                                                                                                                                        CF336199
CF336199.1 GI:33820789
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School of Biological Sciences,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Neoptera; Endopterygota; Diptera; Brachycera; Musco
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Glossina morsitans morsitans
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                   0.7%; So llarity 100.0%; 1 Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Glossina morsitans
/mol_type="mRNA"
/sub_species="morsitans"
/db_xref="taxon:37546"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clone="Tse46d05_p1c"
/tissue_type="adult infected gut"
/clone_lib="Glossina morsitans morsitans adult infected
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       note="country: Zimbabwe; EST
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5.6e+02;
hes 0;
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Muscomorpha;
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REFERENCE
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T52836/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20;
                                                                                                                                                                                                                                                                                                Hillier, L., Lennon, G., Becker, M., Bonaldo, M.F., Chiapelli, B., Chiasee, S., Dietrich, N., DuBuque, T., Favello, A., Gish, W., Hawkins, M., Hultman, M., Kucaba, T., Lacy, M., Le, M., Le, N., Mardis, E., Moore, B., Morris, M., Parsons, J., Prange, C., Rifkin, L., Rohlfing, T., Schellenberg, K., Soares, M.B., Tan, F., Thierry-Meg, J., Trevaskis, E., Underwood, K., Wohldmann, P., Waterston, R., Wilson, R.
                                                                                                          Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. 1 Tel: 314 286 1800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               752836 28 bp mRNA linear EST 06-
ya81b09:s1 Stratagene ovary (#937217) Homo sapiens cDNA clu
IMAGE:68057 3' similar to similar to gb:x53463 GLUTATHIONE
PEROXIDASE-GASTROINTESTINAL (HUMAN), mRNA sequence.
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Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
                                                                                                                                                                                                                                               Generation and analysis of 280,000 human Genome Res. 6 (9), 807-828 (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 28)
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High qality sequence starts: 1 High qality sequence stops: 1 Source: IMAGE Consortium, LLNL This clone is available royal
                                                                   Email: est@watson.wustl.edu
                                                                                          Fax: 314 286 1810
                                                                                                                                                                                Contact: Wilson RK
                                                                                                                                                                                                        8889549
                                                                                                                                                                                                                            97044478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens (human)
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            of Bioscience and Bioinformatics,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Nahm B.H.
Genomics and Genetics Institute,
                                              Insert Size: 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note="Vector: pCR4-TOPO; Site_1: EcoRI; Oligo-capped mRNA
was reverse transcribed and then used for PCR. mRNA was
pepared from Arabidopsis Jasmoinate Carboxyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     methyltransferase overexpression line."
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/lab_host="E.coli DH10B"
/clone_lib="ALJMT-overexpressing transgenic rice
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    organism="Oryza sativa
/mol_type="mRNA"
/cultivar="Nackdong"
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/clone="JMT--06-C18"
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100.0%; Pred. No.
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5.6e+02;
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, MyongJi University
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    available royalty-free
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through LLNL ; contact the IMAGE

Consortium (info@image.llnl.gov)

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JOURNAL COMMENT
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AUTHORS
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AZ654007/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20;
                                                                                                                                                                               Email: ddunn@genetics.utah.edu
Insert Length: 1000 Std Error: 0.00
Plate: 0527 row: 0 column: 20
Seg primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                     Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. Mouse, whole genome scaffolding with paired end reads from 10kb Mouse, whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                               University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AZ654007 28 bp DNA linear GSS 1-
1M0527020R Mouse 10kb plasmid UUGC1M library Mus musculus
clone UUGC1M0527020 R, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (2000)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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801 585 7177
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                          USA
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/dev_stage="49 year old"
/lab host="SOLR cells (Kanamycin resistant)"
/clone_lib="SULR cells (Kanamycin resistant)"
/clone_lib="Stratagene ovary (#937217)"
/note="Organ: ovary; Vector: Bluescript SK; Site 1: EcoRI;
Site 2: XhOI, Cloned unidirectionally. Primer: Oligo dT.
Total ovary tissue, normal, caucasian. Average insert
size: 0.8 kb; Uni-ZAP XR Vector; -5' adaptor sequence: 5'
GAATTCGCACGAG 3' -3' adaptor sequence: 5'
GAATTCGTCACGAG 3' -3' adaptor sequence: 5'
                                                                                                                                                                                                                                                                                                                      Biomedical Polymers Research Bldg.,
                                            /organism="Mus musculus"
/mol_type="genomic DNA"
/strain="C57BL/6J"
          /db_xref="taxon:10090"
/clone="UUGC1M0527020"
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/mol_type="mRNA"
/db_xref="GDB:504314"
/db_xref="taxon:9606"
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100.0%; Pred. No. 5.6e+02;
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ACAAAAAAAAAAAAAAAA 28
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Unpublished (1999)
Contact: MIPS
                                 Similarity
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Location/Qualifiers
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Ottenwaelder, B., Obermaier, B.,
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DKFZp566L2146_r1 566 (synonym: hfkd2) Homo sapiens cDNA clone DKFZp566L2146, mRNA sequence.
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     0.7%; So lilarity 100.0%; E Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pDM12 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                 /organism="Homo sapiens"
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/tissue_type="ridney"
/dev_stage="fetal"
/lab host="X1-2blue"
/clone_lib="566 (synonym: hfkd2)"
/note="Vector: pAMP1; Site_1: NotI; Site_2: Sa
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                              Score 20; Pred. No.
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                              DB 1; Lo
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BX553986/c
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Best Local :
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Lehane, M.J., Aksoy, S., Gibson, W., Kerhornou, A., Berriman, M.,
Hamilton, J., Soares, M.B., Bonaldo, M.F., Lehane, S. and Hall, N.
Adult midgut expressed sequence tags from the tsetse fly Glossina
morsitans morsitans and expression analysis of putative immune
           School of Biological Sciences,
University of Wales,
Bangor LL57 2UW
                                                        Contact: Hall N
Pathogen Sequencing Unit
The Sanger Institute The Wellcome Trust Genome Campus
Hinxton, Cambridge, CB10 ISA, UK
Request for clones, please contact: Mike Lehane
Prof. M.J.Lehane
                                                                                                                                                                   response genes
Genome Biol. 4 (10), R63 (2003)
22881942
                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Hippoboscoidea; Glossinidae; Glossina.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Location/Qualifiers
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ilarity 100.0%;
Conservative
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/mol type="mRNA"
/db xref="taxon:9606"
/clone="DKRZD56411472"
/tissue_type="brain"
/dev_stage="fetal"
/lab_host="X1-Zblue"
/clone_lib="564 (synonym: hfbr2)"
/note="Vector: pAMP1; Site_1: NotI; Site_2: Sa
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5.6e+02;
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BX554037/c
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2660 АСЛАЛАЛАЛАЛАЛАЛАЛА 2679
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BX554037 Glossina consitans cone
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School of Biological Sciences,
University of Wales,
Bangor LL57 2UM
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta;
Neoptera; Endopterygota; Diptera; Brachycera; Muscc
Hippoboscoidea; Glossinidae; Glossina.
                                                                                                                                                                                                                                                                                                                                                                                  Pathogen Sequencing Unit
The Sanger Institute The Wellcome Trust Genome Campus
Hinxton, Cambridge, CB10 1SA, UK
Request for Clones, please contact: Mike Lehane
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Genome Biol. 4 (10),
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s 3' end.
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/db xref="taxon:37546"
/clone="Tpe131902 pic"
/tissue type="adult infected gut"
/clone_Tib="Glossina morsitans mo:
/note="country: Zimbabwe;
I.brucei"
                                                       /db_xref="taxon:37546"
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/tissue_type="adult infected gut"
/clone_Tib="Glossina morsitans morsitans
                                                                                                                                     organism="Glossina morsitans/mol_type="mRNA"

/mol_type="mRNA"

/sub_species="morsitans"
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F.brucei"
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5.6e+02;
                 EST from adult gut
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Best Local Similarity
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                                      1 1M0261I24R Mouse 10kb plasmid U
clone UUGC1M026II24 R, genomic 1
AZ458127
AZ458127.1 GI:10616252
GSS
GSS
Mus musculus (house mouse)
Mus musculus
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Oryza sativa (japonica cultivar-group)
Evaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
Bukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                              AZ458127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeonggi, Korea Tel: 82 31 330 6193 Fax: 82 31 321 6355
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H.,

Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)
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7LEAF--03-N11.bl Rice leaf plasmid cDNA library II (7LEAF) Oryza
8ativa (japonica cultivar-group) cDNA clone 7LEAF--03-N11, mRNA
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                                                                                                                                                                                                                                                                                                                                                                               Conservative
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/clone="71EAF--03-NI1"
/tissue_type="lasf"
/dev_stage="7 days after germination"
/lab_host="E.coli DH10B"
/clone_lib="Rice leaf plasmid cDNA library II (7LEAF)"
/note="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA was capped
with oligoribonuclectides and then used as templates for
RT-PCR."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Oryza sativa (japonica cultivar-group)"
/mol_type="mRNA"
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100.0%; Prr
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Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta
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7LEAF--01-M05.b1
sativa (japonica
                                                                                                                    CF298396
CF298396.1
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Insert Length: 10000 Std Error: (
Plate: 0261 row: I column: 24
Seg primer: CACACAGGAAACAGCTATGACC
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University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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Unpublished (2000)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Class: plasmid ends
High quality sequence stop: 30.
Location/Qualifiers
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Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWH2 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mol_type="genomic DNA'
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/clone="UUGC1M0261124"
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100.0%; Pred. No.
tive 0; Mismatcl
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Streptophyta;

Embryophyta; Tracheophyta;

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CP302456.1 GI:33674217
EST.
Oryza Bativa (japonica cultivar-group)
Oryza Bativa (japonica cultivar-group)
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Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., Le Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H. Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
                                                                                                                                                                                                                                                           Yongin, Kyeonggi, Ko
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                              Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeonggi, Korea
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., L.

Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)
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19 bp mRNA linear EST 15-AUG-2003
7LEAF--07-P22.bl Rice leaf plasmid cDNA library II (7LEAF) Oryza
sativa (japonica cultivar-group) cDNA clone 7LEAF--07-P22, mRNA
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
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Genomics and Genetics Institute, of Bioscience and Bioinformatics,
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llarity 100.0%; Pred. No. 1.8e+03;
Conservative 0; Mismatches 0;
                                                                                                                                                                                                    bhnahm@ggbio.com, bhnahm@bio.myongji.ac.
Location/Qualifiers
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/lab_host="E.coli DH10B"
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/note="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA was cappe
with oligoribonucleotides and then used as templates fc
RT-PCR."
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/cultivar="Nackdong"
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                                                                                                                                              (japonica cultivar-group)"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lee, T.H.,
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Best Local 9
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AL587572 BP Chicken Brain ROS059B11, mRNA sequence.
Gallus gallus (chicken)
Gallus gallus
Gallus gallus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordatae; Galliformes; Phasianidae;
Phasianinae; Gallus.
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AL587572.1
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: MIPS
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Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Wiemann,S.
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Ottenwaelder, B., Obermaier, B.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
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/lab_host="E.coli DH10B"
/clone lib="Rice leaf plasmid cDNA library II (7LEAF)"
/note="Vector: pCR4-TOPO; Site 1: EccRI; mRNA was capped
with oligoribonucleotides and then used as templates for
RT-PCR."
                                                                                                                                                                                                                                                                                                                                                                                                                                   /tissue_type="kidney"
/dev stage="fetal"
/lab_host="X1-2blue"
/clone_lib="566 (synonym: hfkd2)"
/note="Vector: pAMP1, Site_1: NotI; Site_2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /mol_type="mRNA"
/db_xref="taxon:9606"
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                                                                                                            GI:13192606
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hes 0;
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AL587727/c
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AL587727 BP Chicken Brain Library Gallus gallus cDNA ROS061001, mRNA sequence.
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19; Conserv
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EST.
                                                                                                                                                                                                          Unpublished (2001)
Contact: Frazer Murray
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                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae; Gallus.
                                                                                                                                                                                                                                                                                                                  Gallus gallus (chicken)
Gallus gallus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Roslin Institute
Roslin, Midlothian, EH25 9PS, UK
Tel: +44 (0)131 527 4200
Fax: +44 (0)131 440 0434
Email: frazer.murray@bbsrc.ac.uk
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Contact: Frazer Murray
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Gallus gallus"
/mol_type="mRNA"
/db_xref="taxon:9031"
/clone="ROS061D01"
                                                                     ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Gallus gallus"
/mol_type="mRNA"
/db_xref="taxon:9031"
/clone="ROS059B11"
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lab_host="DH108"
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100.0%; Pred. No. 1.8e+03;
tive 0; Mismatches 0;
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Oryza sativa (japonica cultivar-group)
Sukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Nahm B.H.
Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Large-scale Sequencing Analysis Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H., Shin,Y.C., Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H. Large-scale Sequencing Analysis of Rice ESTs
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CF282002.1 GI:33659389
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                                                                                            Similarity
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CAAAAAAAAAAAAAAAA 2
                                    САЛАЛАЛАЛАЛАЛАЛАЛА 2679
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                                                                                                                                                                                                                                                                                                                                                                                                                                              bhnahm@ggbio.com, bhnahm@bio.myongji.ac.
Location/Qualifiers
                                                                                                                                                                                                                                                                /organism="Oryza sativa (japonica cultiv mol type="mRNA" (cultivar="Nackdong" /db_xref="taxon: 39947" /clone="14ETL--09-F01" /tissue_type="leaf" /dev_stage="14 days after germination" /lab_host="E.coli DH10B"
                                                                                                                                                                     /note="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA was capped with oligoribonuclectides and then used as templates for RT-PCR."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /dev_stage="Unknown"
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/clone_lib="BP_Chicken Brain Library"
/note="Vector: pSPORT1; Site_1: Not1; Site_2: Sal1; Cloned unidirectionally. Primer: Oligo dT. 5' adaptor sequence: 5' TCGACCTCGAG 3', 3' adaptor sequence: 5' GCGCCCCTTTTTTTTTTTTTTTTTTTTT 3' Poly A RNA purchased from Clonetech (*6854-1)"
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                                                                                          1.8e+03;
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RESULT 25 CF316662/c

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CF320843/c
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hes 19; Conservative
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                                                                                                                                                                                                                                                                                      20 bp mRNA linear EST 15-AUG HD--11-O01.g1 OsHDAC1-overexpressing transgenic rice plasmid cllbrary (HD) Oryza sativa (japonica cultivar-group) cDNA clone HD--11-O01, mRNA sequence.
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CF316662.1 GI:33688423
                                        1 (bases 1 to 20)

Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., L.

Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)
                                                                                                                                          Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta, Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoidaes; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Nahm B.H.

Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H., Shin,Y.C., Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H. Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
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oryza sativa (japonica cultivar-group)
oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
                           Contact: Nahm
                                                                                                                                                                                                                                                                     CF320843.1 GI:33692604
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
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Location/Qualifiers
      and Genetics
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/dev_stage="proliferated callus on 2N6 media for 2 weeks"
/lab_host=-E_coli DH10B"
/Clone_lib="OsHDAC1-overexpressing transgenic rice plasmid cDNA_Ibrary (HD)"
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/db_xref="taxon:39947"
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100.0%; Pred. No.
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. 1.8e+03;
    GreenGene Biotech
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19; Conserv
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Nouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AZ579178 20 bp DNA linear GSS 13-DEC-20: 1M0363F11F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0363F11 F, genomic survey sequence.
                                                                                                                         High quality sequence stop: 20.
Location/Qualifiers
                                                                                                                                                                                                        Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0363 row: F column: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
1 (bases 1 to 20)
                                                                                                                                                               Class: plasmid ends
                                                                                                                                                                                     Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                   Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                University of Utah Genome University of Utah
                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert
                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished (2000)
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                                                                                                                                                                                                                                                                                                                               Biomedical Polymers Research Bldg.,
                                  /organism="Mus musculus"
/mol_type="genomic DNA"
/strain="C57BL/6J"
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/db_xref="taxon:10090"
/clone="UUGC1M0363F11"
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of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeonggi, Korea Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                           /note="Vector: pCR4-TOPO; Site 1: EcoRI; Callus was treated with ABA(20um) for 1hr. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was derived from rice Histone Deacetylase overexpression
                                                                                                                                                                                                                                                             dev_stage="proliferated callus on 2N6 media for 2 weeks"
lab_host="E.coli DH10B"
/clone_lib="05HDAC1-overexpressing transgenic rice plasmid
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/mol_type="mRNA"
/cultivar≈"Nackdong"
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                      BB
1.8e+03;
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                        6,
                      Length
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SOURCE
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Matches 19; Conserv
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                                                                                                                                 Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0094 row: D column: 20
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
                                                                                                                                                                                                                                                                      University of Utah Genome Center University of Utah Rm. 308, Biomedical Polymers Rese
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Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                 Mouse whole genome scaffolding with paired end reads from 10kb
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Mammalia; Eutheria; Rodentia;
1 (bases 1 to 20)
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                                                                                                       quality sequence stop: 20.
Location/Qualifiers
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801 585 7177
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      /mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC2M0094D20"
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                                                                     organism="Mus musculus"
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100.0%; Prr
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Pred. No.
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RESULT 28 AZ821905

밁 Ş ORIGIN

ACCESSION VERSION

ŒYWORDS

JOURNAL COMMENT

FEATURES

REFERENCE AUTHORS

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AL038582
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DKFZp566F0946 rl 566 (synonym:
DKFZp566F0946, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 21)
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    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: MIPS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ottenwaelder, B., Obermaier, B., Mewes, H.W., Gassenhuber, J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wiemann,S.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /lab_host="E. Coli strain XL10-Gold, T1-resistant, Clone lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: pw1042nv; Purified genomic DNA from musculus C57BL/6J (male) was obtained from the Jackboratory Mouse DNA Resource
                                                                                                                                                                           /lab_host="X1-2blue"
/clone_lib="566 (synonym: hfkd2)"
/note="Vector: pAMP1; Site_1: NotI; Site_2: SalI"
                                                                                                                                                                                                                                                                                                             organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                /clone="DKFZp566F0946"
/tissue_type="kidney"
/dev_stage="fetal"
/lab_host="X1-2blue"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GI:49682163
                                                                             0.7%; Score 19; DB 1; I
100.0%; Pred. No. 1.8e+03;
ative 0; Mismatches 0;
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100.0%; Pred. No.
20
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hfkd2) Homo sapiens cDNA clone
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. 1.8e+03;
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                                                                                                                    Length 21;
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                                                                               Indels
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CF276747/c
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Best Local
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                                                                   Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeonggi, Korea Tel: 82 31 330 6193 Fax: 82 31 321 6355
                                                                                                                                                                            1 (bases 1 to 21)

Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., Lee, T.H., Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)
                                                                                                                                                                                                                                                                  Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                          CF276747
21 bp mRNA linear
14ETL--02-A06.bl Rice etiolated leaf plasmid cDNA
Oryza sativa (japonica cultivar-group) cDNA clone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EST (Ottenwaelder, et al.)
Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 21)
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DKFZp566P1346 rl 566 (synonym:
DKFZp566P1346, mRNA sequence.
                                                                                                                                                           Contact: Nahm B.H
                                                                                                                                                                                                                                                                                                                                                                         CF276747.1 GI:33654133
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ottenwaelder, B., Obermaier, B.,
                                                                                                                                                                                                                                                                                                                                                                                                          mRNA sequence.
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                  bhnahm@ggbio.com, bhnahm@bio.myongji.ac Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mol type="mRNA"
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/clone="DK72p56661346"
/tissue_type="kidney"
/dev_stage="fetal"
/lab_host="X1-2blue"
/clone_lib="566 (synonym: hfkd2)"
/note="Vector: pAMP1; Site_1: NotI;
1. .21
/organism="Oryza sativa
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Pred. No.
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(japonica cultivar-group)"
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library (14ETL)
14ETL--02-A06,
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2661 СААЛАЛАЛАЛАЛАЛАЛА 2679

Mismatches

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В 1.8e+03;

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CF311914/c
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        Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Genomics and Genetics Institute, GreenGene of Bioscience and Bioinformatics, MyongJi U Yongin, Kyeonggi, Korea Tel: 82 31 330 6193 Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H., Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H. Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta, Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CF311914 21 bp mRNA linear EST 15-AUG-2003 ABF--07-G07.gl ABF3-overexpressing transgenic rice plasmid cDNA library (ABF) Oryza sativa (japonica cultivar-group) cDNA clone ABF--07-G07, mRNA sequence.
CF311914 GI:33683675
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Nahm B.H.
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0.7%; Score 19; larity 100.0%; Pred. No. Conservative 0; Mismatc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              bhnahm@ggbio.com,
                                                                                                   /tissue_type="ladf"
/dev stage="14 days after germination"
/lab_host="E.coli DH10B"
/clome llb="ABF3-overexpressing transgenic rice plasmid cDMA llbrary (ABF)"
/note="Vector: pCR4-TOPO; Site_1: EcoRI; Leaf was dried for 2hrs. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was prepared from ABA-responsive element binding transcription factor 3 overexpression line."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA was capped
with oligoribonucleotides and then used as templates for
                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                /db_xref="taxon:39947"
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                                                                                                                                                                                                                                                                                                                                                         organism="Oryza sativa
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/cultivar="Nackdong"
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/dev_stage="14_days_after_germination"
/lab_host="E.coli_DH10B"
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%; Pred. No. 1.8
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                            bhnahm@bio.myongji.ac.kr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 6; Le
1.8e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MyongJi University
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CF328832/c
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                 FEATURES
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Best Local Similarity
Matches 19; Conserv
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source
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                                                                                                                                                                                                                     Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
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                                                                                           Genomics and Genetics Institute, Cof Bioscience and Bioinformatics,
                                                                                                                                 Contact: Nahm B.H.
                                                                                                                                                 1 (bases 1 to 22)

Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Le

Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)
                                                                                                                                                                                                                                                                                                                       EST.
                                                                                                                                                                                                                                                                                                                                   sequence.
CF328832
CF328832.1 GI:33805905
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1 (bases 1 to 22)
Koehrer,K., Beyer,A., Mewes,H.W.,
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22 bp mRNA linear EST 04-S DKFZp5566L123 r1 566 (synonym: hfkd2) Homo sapiens cDNA clone DKFZp566L123, mRNA sequence.
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Location/Qualifiers
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82 31 330 6193
82 31 321 6355
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            bhnahm@ggbio.com, bhn
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Homo sapiens"

/mol type="mRNA"

/db_xref="taxon:9606"

/clone="DKFZp5661123"

/tissue_type="kidney"

/dev_stage="fetal"

/lab host="X1-2blue"

/clone_lib="566 (synonym: hfkd2)"

/note="Vector: pAMP1; Site_1: Not1; Site_2: Sal1"
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%; Pred. No. 1.8
0; Mismatches
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callus plasmid cDNA library (NACL) Oryza
tivar-group) cDNA clone NACL--03-020 מייאים
                          bhnahm@bio.myongji.ac.kr.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.8e+03;
thes 0;
                                                                                          GreenGene Biotech Inc.; Division, MyongJi University
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                                                                                                                                                                                             Lee, T.H.,
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0005 row: K column: 17
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert B. University of Utah University of Utah
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Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
1 (bases 1 to 22)
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Location/Qualifiers
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/mol type="mRNA"
/cultīvar="Nackdong"
/db_xref="taxon:39947"
/clone="NACL--03-020"
/tissue_type="callus"
/dev_stage="proliferated callus on 2N6 media for 30 days"
/dev_stage="proliferated callus on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       USA
(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4
                                                                                                                         Biomedical
                                                                                                                                                                                                                                                                                                                                                                                     /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                 sex="Male"
                                                                                                                                                                                                                                                                                                                                db_xref="taxon:10090"
clone="UUGC1M0005K17"
                                                                                                                                                                                                                                                                                                                                                                                                                                                            organism="Mus musculus"
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Pred. No.
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mid UUGC1M library Mus musculus genomic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 High quality sequence stop: 22.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0346 row: A column: 10
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.

Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 22)

Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AZ505769 22 bp DNA linear GSS 05-OCT-20
1M0346A10R Mouse 10kb plasmid UUGCIM library Mus musculus genomic
clone UUGC1M0346A10 R, genomic survey sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (2000)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   USA
/lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone lib="Wouse 10kb plasmid UUGCIM library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C578L/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Biomedical Polymers Research Bldg., 20 S. 2030 E.,
                                                                                                                                                                                                                                                                                                               /db_xref="taxon:10090"
/clone="UUGC1M0346A10"
                                                                                                                                                                                                                                                                                                                                                                        /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                        organism="Mus musculus"
                                                                                                                                                                                                                                                                                     'sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.7%;
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Pred. No.
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1.8e+03;
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REFERENCE AUTHORS

COMMENT

JOURNAL

FEATURES

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RESULT 36 AZ505769 LOCUS

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Matches

ACCESSION VERSION

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                                                                                                                 Query Match
Best Local
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Best Local :
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                                          2661 САЛАЛАЛАЛАЛАЛАЛАЛА 2679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2661 САРАРАРАРАРАРАРА 2679
20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Roslin, Midlothian, BH25 9PS, UNITED KINGDOM
Single pass sequencing. Bases called and trimmed with phred
v0.020425.c. Vector identified by cross match with the -minscore 20
and -minmatch 12 options. Vector:pBlueScriptII(KS+) R. Site 1:
ECORI R. Site 2: NotI Description: Normalised library constructed
from pooled tissue from day 30 placentas. Clones available from UK
Centre for Functional Genomics in Farm Animals, Roslin Institite,
Roslin, Midlothian, UK, EH25 9PS, www.arkgenomics.org.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Anderson, S.I., Finlayson, H.A. and Archibald, A.L. Development of CDNA and EST resources for studying reproduction and embryo development in pigs and cattle Unpublished (2004)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            שמאמשת Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae; Sus.

1 (bases 1 to 23)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AJ666332 CSEQRAN09 Sus scrofa cDNA clone C0000033_H19, mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Anderson SI
Genomics and Bioinformatics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sus scrofa (pig)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AJ666332.1
EST.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gi | 4732114 | gb | AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                              /mol_type="mRNA"
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/db_xref="taxon:9823"
/db_xref="taxon:9823"
/clone="C0000003_H19"
/tissue_type="placenta"
/clone_Tib="CSEQRAN09"
/clone_Tib="CSEQRAN09"
/note="Vector: pBlueScriptII(KS+); Site_1: EcoRI; Site_2:
/note="Vector: pBlueScriptII(KS+); Normalised library
constructed from pooled tissue from day 30 placentas."
                                                                                                                                                                                                                                                                                                                                                                                                                                                   ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                      organism="Sus scrofa"
                                                                               100.0%; Fi
                                                                                                            0.7%; Score 19; DB 1;
00.0%; Pred. No. 1.8e+0
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                                                                                         1.8e+03;
hes 0;
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hes 0;
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                                                                                                                                   Length 23;
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SOURCE

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AUTHORS
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CF318266/c
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AUTHORS
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AL048745
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VERSION
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Best Local Similarity
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                                                                                                                                                                                                                                           1 (bases 1 to 23)

Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., L.

Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                 Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
                                                                                                                                Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                 Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                                                               Contact: Nahm B.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CF318266 RNA linear EHD--08-F07.bl OsHDAC1-overexpressing transgenic rice library (HD) Oryza sativa (japonica cultivar-group) c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4 CAAAAAAAAAAAAAAAA 22
                                                                                                                                                                                                                                                                                                                                              Ehrhartoideae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          CF318266.1 GI:33690027
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HD--08-F07, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 (bases 1 to 23)
Koehrer, K., Beyer, A.,
EST (Koehrer, et al.)
Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AL048745 23 bp mRNA DKFZp566K213 rl 566 (synonym: hfkd2) Hon DKFZp566K213, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: MIPS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo
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                                                                                          bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr
Location/Qualifiers
               /organism="Oryza sativa (japonica cultivar-group)"
/mol_type="mRNA"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone_lib="566 (synonym: hfkd2)"
/note="Vector: pawP1; Site_1: NotI; Site_2: SalI"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /tissue_type="kidney"
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/clone="DKFZp566K213"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GI:4727816
_xref="taxon:39947"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.7%; Score 19; DB 100.0%; Pred. No. 1.8 tive 0; Mismatches
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thes 0;
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Homo sapiens cDNA clone
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                                                                                                                                                                                                                                                                                                  Lee, T.H., Shin, Y.C.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            e plasmid cl
cDNA clone
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JOURNAL COMMENT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: ddurn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0033 row: O column: 04
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert B. Weiss University of Utah Genome University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokee,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 23)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1M0033004F Mouse 10kb plasmid UUGC1M library McClone UUGC1M0033004 F, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                          High quality sequence stop: 23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       plasmid inserts
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            . 308,
USA
Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were
                                                                                                                                        /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/Clone lib="Mouse 10kb plasmid UUGCIM library"/note="Vector: PWD42nv; Purified genomic DNA from M.musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                          /organism="Mus musculus"
/mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone lib="OsHDAC1-overexpressing transgenic rice plasmid cDNA library (HD)"
/note="Vector: pCR4-TOPO; Site_1: EcoRI; Callus was
                                                                                                                                                                                                                                 /clone="UUGC1M0033004"
/sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  treated with ABA(20um) for 1hr. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was derived from rice Histone Deacetylase overexpression
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/tissue_type="callus"
/dev_stage="proliferated callus on
/lab_host="E.coli DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GI:10362861
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Pred. No.
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CF326993
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TITLE
JOURNAL
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Best Local Similarity
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                                                                             CF326993 24 bp mRNA linear EST 10 NACL--01-E17.gl Rice callus plasmid cDNA library (NACL) 01 sativa (japonica cultivar-group) cDNA clone NACL--01-E17,
                                                                                                                                                                                          4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 (bases 1 to 24)
Koehrer,K., Beyer,A., Mewes,H.W., Gassenhuber,J. and Wiemann,S.
EST (Koehrer, et al.)
Unpublished (1999)
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EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24 b
DKFZD566M233 rl 566 (synonym:
DKFZD566M233, mRNA sequence
AL048765
EST.
Oryza sativa (japonica cultivar-group)
                                                     sequence.
CF326993
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                   CF326993.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CAAAAAAAAAAAAAAAAA 21
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                                                                                                                                                                                                                                                                                                                         /tissue_type="kidney"
/dev_stage="fetal"
/lab_host="X1-2blue"
/clone_lib="566 (synonym: hfkd2)"
/note="Vector: pAMP1; Site_1: Not1;
                                                                                                                                                                                                                                                                                                                                                                                                                        organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ocation/Qualifiers
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                                   GI:33802241
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100.0%; Pr
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100.0%; Pred. No. 1.8e+03;
ive 0; Mismatches 0;
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1.8e+03;
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Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., Lee, T.H., Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H. Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0173 row: J column: 14
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                    Contact: Robert B. Weiss
University of Utah Genome
University of Utah
                                                                                                                                                                                                                                                                                                                                                                               1 (bases 1 to 24)
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B.
Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Peder
Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A.,
Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AZ404871 24 bp DNA linear GSS 03-OCT-20 1M0173J14R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0173J14 R, genomic survey sequence.
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Yongin, Kyeenggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oryza sativa (japonica cultivar-group)
Gukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
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                                                                                                              Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                      Unpublished (2000)
                                                                                                                                                                                                                                                                                                                               plasmid inserts
                                                                                                                                                                                                                                                                                                                                                         Mouse whole genome scaffolding with paired end reads from 10kb
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Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus
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                                                                                                                                                                                             Biomedical Polymers Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA was capped with oligoribonucleotides and then used as templates for RT-pCR "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /tissue_type="callus"
/dev stage="proliferated callus on 2N6 media for 30 days"
/lab_host="E.coll DH10B"
/clome_lib="Rice callus plasmid cDNA library (NACL)"
/clome_lib="Rice callus plasmid cDNA library (NACL)"
/clome_lib="Vector: pCR4-TOPO; Site_l: ECRN; mRNA was capped
/inte="Vector: pCR4-TOPO; Site_l: ECRN; mRNA was capped
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Oryza sativa
/mol_type="mRNA"
/cultivar="Nackdong"
/db_xref="taxon:39947"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
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100.0%; Pred. No.
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                                                                                                                                                                                                                                                      Center
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.8e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Duval, B., Hamil, C.,
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AZ786257
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Best Local Similarity
Matches 19; Conserv
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                                                                                                                                                                                                                                                                                                                                                     Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. Mouse whole genome scaffolding with paired end reads from 10kb
                                 Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0031 row: H column: 11
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                         Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                         University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                               plasmid inserts
Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 24)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AZ786257 24 bp DNA linear GSS 16 2M0031H11R Mouse 10kb plasmid UUGC1M library Mus musculus clone UUGC2M0031H11 R, genomic survey sequence.
AZ786257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4 CAAAAAAAAAAAAAAAAA 22
               Class: plasmid ends
                                                                                                                                                                                                                                                                                            Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AZ786257.1 GI:12923835
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus (house mouse)
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Location/Qualifiers
                                                                                                                                                                             . 308,
USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      / Tab. host="E. Coli strain XL10-Gold, T1-resistant, F-"
/ (clone lib="Mouse 10kb plasmid UUGCIM library"
/ (note="Vector: PMD42nv; Purified genomic DNA from M.
musculus C57EL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polymucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD42 (gil4732114|gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
chemically-competent E. coli XL10-Gold (Stratagene) cells
and selected for ampicillin resistance."
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/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0173J14"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       'sex="Male"
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                                                                                                                                                                                                                                                                                            Weiss
                                                                                  Std Error: 0.00
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                                                                                                                                                                                                          S. 2030 E.,
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COMMENT
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AUTHORS
TITLE
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KEYWORDS
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AL587718/c
                                                                                                          FEATURES
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                                                                                                                                                                     Murray, F.
BP Chicken Brain Library
Unpublished (2001)
                                                                                                                                                                                                                                                                                                                 Contact: Frazer Murray
Dept. Genomics and Bioinformatics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gallus gallus (chicken)
Gallus gallus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AL587718 25 bp mRNA linear EST 02
AL587718 BP Chicken Brain Library Gallus gallus cDNA clone
ROS061C02, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae; Gallus.
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AL587718.1 GI:13192752
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        (bases 1 to 25)
                                                                                                                              primer: M13F
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Individual DNA Resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pMN42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
/organism="Gallus gallus"
/mol_type="mRNA"
/db_xref="taxon:9031"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone_lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                     ocation/Qualifiers,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /mol_type="genomic DNA'
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 clone="UUGC2M0031H11"
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L00.0%; Pred. No. 1.8e+(
ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             o;
                                                                                                                                                                     Poly A RNA purchased from Clonetech
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Length 24; Indels

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N27663/c
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PUBMED
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JOURNAL
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                                                                                                                                                                                                                                                                                   source
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                                                                                                                                                                                                                                                                                                                    High quality sequence starts: 1
High quality sequence stops: 1
Source: IMAGE Consortium, LLNL
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.lln1.gov) for further information.
Trace considered overall poor quality
Seq primer: m13 -40 forward
High quality sequence stop: 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hillier, L., Lennon, G., Becker, M., Bonaldo, M.F., Chiapelli, B., Chissoe, S., Dietrich, N., DuBuque, T., Favello, A., Gish, W., Hawkins, M., Hultman, M., Kucaba, T., Lacy, M., Le, M., Le, N., Mardis, E., Moore, B., Morris, M., Parsons, J., Prange, C., Rifkin, L., Rohlfing, T., Schellenberg, K., Soares, M.B., Tan, F., Thierry-Meg, J., Trevaskis, E., Underwood, K., Wohldmann, P., Waterston, R., Wilson, R. and Marra, M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (HUMAN);, mRNA sequence.
N27663
N27663.1 GI:1142144
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yw50g06.sl Weizmann Olfactory Epithelium Homo sapiens cDNA cl
IMAGE:255706 3' similar to gb:J05032 ASPARTYL-TRNA SYNTHETASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Genome Res.
97044478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Generation and analysis of 280,000 human expressed sequence tags Genome Res. 6 (9), 807-828 (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1. (bases 1 to 25)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens (human)
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                                                                                                                                                     /organism="Homo sapiens"
/mol type="mRNA"
/db xxef="GDB:3866115"
/db xxef="taxon:9606"
/clone="IMAGE:255706"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /dev stage="Unknown"
/lab_host="DH10B"
/clone lib="BP Chicken Brain Library"
/clone lib="BP Chicken Brain Library"
/cloned:
/note="Vector: pSPORT1; Site 1: Not1; Site 2: Sal1; Cloned unidirectionally. Primer: Oligo dT. 5' adaptor sequence: 5' TCGACCTCGAG 3'; 3' adaptor sequence: 5'
GCGGCCGCTTTTTTTTTTTTTTTTTTT 3' Poly A RNA purchased from
     /dev stage="35 year old"
/lab_host="SOLR cells (kanamycin resistant)"
/clone_lib="Weizmann Olfactory Epithelium"
/note="Organ: nose; Vector: pBluescript SK-;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Clonetech (*6854-1)"
                                                                                                       tissue_type="olfactory epithelium"
                                                                                                                                      'sex="Female"
                                                                                                                                                                                                                                                                                                     ocation/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone="ROS061C02"
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100.0%; Pr/
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Louis, MO 63108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 25;
SK-; Site_1:
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SOURCE

COMMENT

Query Match Best Local S Matches 19

l Similarity 19; Conserv

Conservative

100.0%; E ative 0;

0.7%; Score 19; DB 9; 100.0%; Pred. No. 1.8e+(

Mismatches

1.8e+03; hes 0;

Indels

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Gaps

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Length 25;

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Beat

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REFERENCE
AUTHORS
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CG726337/c
LOCUS
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VERSION
ORIGIN
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JOURNAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Stanford University
855 California Ave,
Tel: 650 723 2227
Fax: 650 725 8221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22 CAAAAAAAAAAAAAAAA 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Reverse complemented post-ligation plate: 111908 row: E column: 12 Class: transposon-tagged.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (2001)
Contact: Walbot V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CG726337 25 bp DNA linear 1119089E12.2EL_yl 1119 - RescueMu Grid AA Zea mays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: walbot@stanford.edu
Possible ligation site of ends cut
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Department of Biological Sciences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Maize genomic sequences found using engineered RescueMu transposon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Walbot, V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               clade; Panicoideae; Andropogoneae;
1 (bases 1 to 25)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GSS.
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CG726337.1
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           /clone lib="1119 - RescueMu Grid AA"
/clone lib="Organ: leaf; Vector: RescueMu (engineered from pBlueScript backbone); Site 1: BamHI; Site 2: BglII;
RescueMu is a 4.9 kb, modified maize Mu transposon designed to allow plasmid rescue from total genomic DNA.
Mu elements insert preferentially into transcription units. For more information on RescueMu, go to the web site 'www.zmdb.lastate.edu' and follow the links for 'RescueMu.' Grid AA was grown at UC San Diego in 2002. DNA was extracted from leaf strips, double digested using BamHI and BglII, and ligated to form circular plasmids. DH10B cells were transformed and then screened on LB plates with ampicillin."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EcoRI; Site 2: XhoI; Cloned unidirectionally. Primer Oligo dT. Olfactory epithelium, normal Average inser size: 0.8 kb; Uni-ZAP XR Vector. Library constructed! Walker, D. Lancet, Welzmann Institute of Science. ~5' adaptor sequence: 5' GAATTCGGCACGAG 3' ~3' adaptor sequence: 5' CTCGAGTTTTTTTTTTTTTTTTT 3'"
                                                                                                                                                                                                                                                                                                   /tissue_type="leaf"
/dev_stage="adult"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                   /mol_type="genomic_DNA"
/cultivar="mixed_background_W23/A188/B73/K55"
/db_xref="taxon:4577"
                                                                                                                                                                                                                                                                                                                                                                                                                                     organism="Zea mays"
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RESULT 49
CN545723/c
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VERSION
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VERSION
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ORGANISM
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CF280688/c
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Best Local S
Matches 19
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                                      AUTHORS
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                                                                                                                                                                                                                                                                                                                                                       21
                                                                                                                                                                                                                CN545723
EST 17667 Ripe Grape
clone B3CS00RL005C03
                                                                                                                                                                                                                                                                                                                                                                                                                             l Similarity
19; Conserv
1 (bases 1 to 26)
Abbal.P., Agasse.A., Ageorges.A., Atanassova.R., Barrieu.F.,
Couture,C., Dedaldechamp.F., Delrot.S., Glissant.D., Grimple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23
                                                               Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eurosids; Vitaceae; Vitis.
                                                                                                                         Vitis vinifera
                                                                                                                                                                               CN545723
CN545723.1 GI:46910348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Nahm B.H.

Contact: Nahm B.H.

Genomics and Genetics Institute, GreenGene Biotech Inc.,
of Bioscience and Bioinformatics, MyongJi University

Yongin, Kyeonggi, Korea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 (bases 1 to 26)
Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., Lee, T.H., Shin, Y.C., Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
                                                                                                                                                Vitis vinifera
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CF280688 26 bp mRNA linear EST 14-AUG-2003 (14ETL--07-H09.gl Rice etiolated leaf plasmid cDNA library (14ETL) CT28 sativa (japonica cultivar-group) cDNA clone 14ETL--07-H09,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Yongin, Kyeonggi, Ko
Tel: 82 31 330 6193
Fax: 82 31 321 6355
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larity 100.0%; Pred. No. 1.8e+03;
Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 bhnahm@ggbio.com,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="Vector: pCR4-TOPO;
with oligoribonucleotides
RT-pCR."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /tissue_type="leaf"
/dev_stage="14 days after germination"
/lab_host="E.coli DH10B"
/clone_lib="Rice etiolated leaf plasmid cDNA library
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Oryza sativa
/mol_type="mRNA"
/cultivar="Nackdong"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /db_xref="taxon:39947"
/clone="14ETL--07-H09"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Site 1: EcoRI; mRNA was capped and then used as templates for
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                                                                                   eudicots;
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       Grimplet, J.,
                                                                                                  Tracheophyta;
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7LEAF--02-P02.b1 F
sativa (japonica (
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., L.

Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
                                                                                                                                                                                                                                                   Yongin, Kyeonggi, K
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                     Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                                                                                                                                                                                    Contact: Nahm B.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eĥrhartoideae; Oryzeae; Oryza.
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Generation of Expressed Sequence Tag from Grape Berry
or seeds) at Various Developmental Stages
Unpublished (2002)
Contact: Hamdi S.
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Fax: 00-33-(0)5-57-12-25-48
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Universite de Bordeaux I, Institut National
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71, Avenue 1
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                                                                                                                                                                                                                                bhnahm@ggbio.com,
/tissue_type="leaf"
/dev_stage="7 days after germination"
/lab_host="B.coli DH10B"
/clone_lib="Rice leaf plasmid cDNA library
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/mol_type="mRNA"
/cultivar="Nackdong"
                                                                                    db_xref="taxon:39947"
clone="7LEAF--02-P02"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /dev stage="ripening stage"
/clone_lib="Ripe Grape Skin Triplex2 Library"
/note="Organ: Fruit skin; Vector: Lambda TriplEx2; Site_1:
SfiIA; Site_2: SfiIB; Oriented library"
                                                                                                                                                                                                             ocation/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Vitis vinifera"
/mol type="mRNA"
/cultivar="Cabernet Sauvignon"
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00.0%; Pred. No. 1.8e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                         bhnahm@bio.myongji.ac.kr
                                                                                                                                                              (japonica cultivar-group)
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Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., L. Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H. Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
CF310745 27 bp mRNA linear EST 15-AUG-2003
ABF--05-J07.gl ABF3-overexpressing transgenic rice plasmid cDNA
library (ABF) Oryza sativa (japonica cultivar-group) cDNA clone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Yongin, Kyeonggi, Ko
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Genomics and Genetics Institute, GreenGene Biotech Inc.; of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeonggi, Korea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
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CF310560
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Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Nahm B.H.
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                       note="Vector: pCR4-TOPO; Site_1: EcoRI; Leaf was dried for 2hrs. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was prepared from ABA-responsive element binding transcription factor 3 overexpression
                                                                                                                                                                                                                                                                                                                                                                                                                                        /clone lib="ABF3-overexpressing transgenic rice plasmid
cDNA library (ABF) "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /dev_stage="14 days after germination"
/lab_host="E.coli DH10B"
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/clone="ABF--05-E14"
/tissue_type="leaf"
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/cultivar="Nackdong"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         organism="Oryza sativa (japonica cultivar-group)"
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AUTHORS
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CF311022/c
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AUTHORS
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Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta;
                                     Contact: Nahm B.H.

Genomics and Genetics Institute, Greendene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                              1 (bases 1 to 27)

Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H.,

Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)
                                                                                                                                                                                                              Oryza sativa (japonica cultivar-group) oryza sativa (japonica cultivar-group) oryza sativa (japonica cultivar-group) Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae; Oryzeae; Oryza.
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Contact: Nahm B.H.

Genomics and Genetics Institute, GreenGene Biotech II

Genomics and Bioinformatics, MyongJi University
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H.,
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
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CF310745
CF310745.1 GI:33682506
                                   of Bioscience and Bioinformatics, Yongin, Kyeonggi, Korea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
Email: bhnahm@ggbio.com,
                                                                                                                                                                                                                                                                                                                                      CF311022.1 GI:33682783
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            Kyeonggi, Ko
31 330 6193
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/mol type="mRNA"
/cultivar="Nackdong"
/db xref="taxon:39947"
/clone="ABF--05-J07"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /dev_stage="14 days after germination"
/lab_host="S.coll DH10B"
/clome_lib="ABP3-overexpressing transgenic rice plasmid cDNA library (ABP)"
/note="Vector: pCR4-TOPO; Site_1: EcoRI; Leaf was dried for Drs. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was prepared from ABA-responsive element binding transcription factor 3 overexpression line."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ocation/Qualifiers
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Abbal, P., Agasse, A., Ageorges, A., Atanassova, R., Barrieu, F., Couture, C., Dedailechamp, F., Detot, S., Glissant, D., Grimplet, J., Hamdi, S., Romieu, C. and Terrier, N.
Generation of Expressed Sequence Tag from Grape Berry (skin, pulp or seeds) at Various Developmental Stages
Unpublished (2002)
                                                                                                                                                                                                                                                            Seq primer: T7
                                                                                                                                                                                                                                                                              Email: s.hamdi@bordeaux.inra.fr
                                                                                                                                                                                                                                                                                                   Tel: 00-33-(0)5-57-12-25-50 Fax: 00-33-(0)5-57-12-25-48
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71, Avenue Edouard Bourleaux,
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Vitis vinifera
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CN545777.1 GI:46910402
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                                                                                                                                                                                                                                                                                                                                                                                                            Universite de Bordeaux I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Hamdi S.
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      /dev_stage="ripening stage"
/clone_lib="Ripe Grape Skin Triplex2 Library"
/note="Organ: Fruit skin; Vector: Lambda TriplEx2; Site_1:
SfilA; Site_2: SfilB; Oriented library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Oryza sativa (
/mol type="mRNA"
/cultivar="Nackdong"
/db_xref="taxon:39947"
/clone="ABF--06-B07"
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/lab_host="E.coli DH10B"
/clone lib="ABF3-overexpressing transgenic rice plasmid cDNA library (ABB)"
/note="Vector: pCR4-TOPO; Site_1: EcoRI; Leaf was dried for 2hrs. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was prepared from ABA-responsive element binding transcription factor 3 overexpression
                                                                                                                                       /organism="Vitis vinifera"
/mol_type="mRNA"
/cultivar="Cabernet Sauvignon"
                                                                                                  /db_xref="taxon:29760"
/clone="B3CS00RL005G11"
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                   Homo sapiens (human)
                                                                                                 N89936 27 bp mRNA linear zb23e12.s1 Soares fetal lung NDHL19W Homo sapiens IMAGE:302926 3' similar to gb:X59066 ATP SYNTHASE MITOCHONDRIAL PRECURSOR (HUMAN);, mRNA sequence.
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                       N89936.1 GI:1443263
                                                                                          N89936
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UMR 619 - Equipe Biologie de la Vigne
Universite de Bordeaux I, Institut National
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
rosids; Vitaceae; Vitis.
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19; Conservative
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/clone_lib="Ripe Grape Skin Triplex2 Library"
/note="Organ: Fruit skin; Vector: Lambda Trip:
SfiIA; Site_2: SfiIB; Oriented library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Vitis vinifera"
/mol type="mRNA"
/cultivar="Cabernet Sauvignon"
/db xref="taxon:29760"
/clone="B3CS00RL002H10"
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T. brucei sheared genomic DNA
genomic survey sequence.
Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and
                                                             Hall, N., Bowman, S., Lennard, N.J., Doggett, J., Atkin, R., Chillingworth, C., Ormond, D., Harris, B., El-Sayed, N., Hou, Melville, S.E., Rajandream, M.A. and Barrell, B.G.
                                                                                                                                                                                                       GSS.
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High quality sequence stop: 8.
Location/Qualifiers
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The WashU-Merck EST Project
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Hiller,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M.,
Holman,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M
Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F.,
Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P. and
                                                                                                                                                                                                                                     AL498287
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: es@watson.wustl.edu
Email: es@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
This clone is available royalty-free through LLNL; contact the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tel: 314 286 1800
Fax: 314 286 1810
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (1995)
Contact: Wilson RK
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                                                                                                                                                Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
                                                                                                                                                                  Trypanosoma brucei
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                                                     Direct Submission
                                                                                                                                                                                    Trypanosoma brucei
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                                                                                                                 (bases 1 to 27)
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/db_xref="GDB:1247858"
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100.0%; Pred. No.
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                             2661 САЛАЛАЛАЛАЛАЛАЛА 2679
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: neleayed@tigr.org
Details of T. brucei sequencing at the Sanger Centre
at http://www.sanger.ac.uk/projects/T_brucei/.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      to give a tight size distribution (
4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned population of Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically sheared
                                                                                                                                                                                                                                                                                                                     Ingolstaedter Landstr.1, D-85764 Neuherberg, Germany.
Location/Qualifiers
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Mammalia; Eutheria; Primates;
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                                                                                                                                              //clone="DKFZp564H2264"
/tissue_type="brain"
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/lab_host="X1-2blue"
/clone_lib="564 (synonym: hf
/note="Vector: pAMP1; Site_1
                                                                                                                                                                                                                                               organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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/clone="386g03"
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|mol_type="genomic DNA"
|strain="TREU927"
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hfbr2) Homo sapiens cDNA clone
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~hes 0;
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RESULT 59

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Best Local (
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Unpublished (1999)
                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 28)

Bloecker, H., Boecher, M., Brandt, P., Mewes, H.W., Gassenhuber, J. and
                                                                                                                                                                                                                                                                                                                                                                                                                          DKFZp564F027 sl 564 (synonym: DKFZp564F027, mRNA sequence.
                                                                                                                                                                      Ingolstaedter Landstr.1, D-8
                                                                                                                                                                                                                                                                      Wiemann,S
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Unpublished (1999)
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AL037270.1 GI:49681975
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1 (bases 1 to 28)
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Homo sapiens
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CAAAAAAAAAAAAAAAA 2679
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism="Homo sapiens"
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/clone_lib="564 (synonym: hfbr2)"
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/clone_lib="564 (synonym: hfbr2)"
/note="Vector: pAMP1; Site_1: NotI; Site_2: SalI"
                                                                                                                         organism="Homo sapiens"
|mol_type="mRNA"
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onym: hfbr2)
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AL587582/c
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AL039180
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JOURNAL
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Matches 1
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Best Local
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         Murray, F.

BP Chicken Brain Library
Unpublished (2001)
Contact: Frazer Murray
Dept. Genomics and Bioinformatics
Roslin Institute
                                                                                               Gallus gallus (chicken)
Gallus gallus
Gallus gallus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Galliformes; Phasianidae;
Phasianinae; Gallus.
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Roslin, Midlothian,
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AL587582.1 GI:13192616
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Unpublished (1999)
Contact: MIPS
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 28)
Bloecker.H., Boecher,M., Brandt,P., Mewes,H.W., Gasse
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DKFZp5660204 rl 566 (synonym:
DKFZp5660204, mRNA sequence.
AL039180
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                                                                                           (bases 1 to 28)
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                                                                                                                                                                                                                                                                                                                                                                                                                       /tissue_type="kidney"
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/lab host="x1-zblue"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RNA linear
Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                       Length 28
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ORGANISM
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BX554747/c
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VERSION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Glossina morsitans morsitans
Glossina morsitans morsitans
Glossina morsitans morsitans
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Hippoboscoidea; Clossinidae; Glossina.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19;
                                                                                                                                                                   Prof. M.J.Lehane Coll. School of Biological Sciences, University of Wales, Bangor LLS7 20W All clones with suffix old are
                                                                                                                                                                                                                                                         Contact: Hall N
Pathogen Sequencing Unit
The Sanger Institute The Wellcome Trust Genome Campus
Hinxton, Cambridge, CB10 1SA, UK
Request for Clones, please contact: Mike Lehane
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lehane,M.J., Aksoy,S., Gibson,W., Kerhornou,A., Berriman,M., Hamilton,J., Soares,M.B., Bonaldo,M.F., Lehane,S. and Hall,N. Adult midgut expressed sequence tags from the tsetse fly Glossina
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28 bp mRNA linear EST 10-OCT-201 MRN554747 Glossina morsitans morsitans adult infected gut Glossina morsitans morsitans morsitans morsitans morsitans morsitans morsitans morsitans clone Tse17c01_p1c, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tel: +44 (0)131 527 4200
Fax: +44 (0)131 440 0434
Email: frazer:murray@bbsrc.ac.uk
GCGGCCGCTTTTTTTTTTTTTTTT 3' Poly A RNA purchased from Clonetech
                                                                                                                                                                                                                                                                                                                                                                                                  response genes
Genome Biol. 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BX554747.1 GI:33378810
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                                                                                                                                                                                                                                                                                                                                                                                                                                     morsitans morsitans and expression analysis of putative immune
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                                                                                                                                  clones with suffix qlc are reverse
of the cDNA all plc reads are from
3' end.
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/note="Vector: pSPORTI; Site 1: Not1; Site 2: Sall; Cloned unidirectionally. Primer: OlTgo dT. 5' adaptor sequence: 5' TCGACCTCGAG 3', 3' adaptor sequence: 5' GCGGCCGCTTTTTTTTTTTTTTTT 3' Poly A RNA purchased from Clonetech (*6854-1)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Gallus gallus"
/mol_type="mRNA"
/db_xref="taxon:9031"
                   /organism="Glossina morsitans
/mol_type="mRNA"
/sub_speciess"morsitans"
/db_xref="taxon:37546"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /dev_stage="Unknown"
/lab_host="DH108"
                                                                                                                 location/Qualifiers
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100.0%; Pr
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; Pred. No. 1.8e+03;
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CN545659/c
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AUTHORS
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CN545498/c
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KEYWORDS
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CN545659 28 bp mRNA EST 17603 Ripe Grape Skin Triplex2 Library clone B3CS00RL004E11 3', mRNA sequence.
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Query Match
Best Local Similarity
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                                     2661 САЛАЛАЛАЛАЛАЛАЛАЛ 2679
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28 CAAAAAAAAAAAAAAAAAA 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Hamdi S.
UMR 619 - Equipe Biologie de la Vigne
Universite de Bordeaux I, Institut National
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Abbal, P., Agasse, A., Ageorges, A., Atanassova, R., Coutture, C., Dedaldechamp, F., Delrot, S., Glissant, Hamdi, S., Romieu, C. and Terrier, M. Generation of Expressed Sequence Tag from Grape For seeds) at Various Developmental Stages Unpublished (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Viridiplantae; Stro
Spermatophyta; Magnoliophyta;
rosids; Vitaceae; Vitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CN545498 28 bp mRNA linear EST 30-APR-:
EST 17442 Green Grape Skin Triplex2 Library Vitis vinifera cDNA
clone B3CS00GL005G11 3', mRNA sequence.
CN545498
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Vitis vinifera
                                                                                                                                                                                                                                                                                                                                                                                                 Email: s.hamdi@bordeaux.inra
Seq primer: T7.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Tel: 00-33-(0)5-57-12-25-50 Fax: 00-33-(0)5-57-12-25-48
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                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /tissue_type="adult infected gut"
/clone_lib="Glossina morsitans morsitans adult infected
gut"
                                                                                                                                                                        /dev stage="green stage"
/clome_lib="Green Grape Skin Triplex2 Library"
/note="Organ: Fruit skin; Vector: Lambda TriplEx2; Site_1:
SfiIA; Site_2: SfiIB; Oriented library"
                                                                                                                                                                                                                                                                                                              /organism="Vitis vinifera"
/mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
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                                                                                                                                                                                                                                                         db_xref="taxon:29760"
clone="B3CS00GL005G11"
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100.0%; Pred. No.
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Pred. No.
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hes 0;
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,D., Grimplet, J.,
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linear EST 30-APR Vitis vinifera cDNA

EST 30-APR-2004

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JOURNAL COMMENT
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AUTHORS
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AZ481286/c
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                                                                                                                                                                                                Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                      University of Utah Genome Center University of Utah
                                                                                                                               Contact: Robert B.
                                                                                                                                               Unpublished (2000)
                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
1 (bases 1 to 28)
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AZ481286
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Tel: 0
                                                                                                                                                                                Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                            Mus musculus
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Abbal, P., Agasse, A., Ageorges, A., Atanassova, R., Barrieu, F.,
Abbal, P., Agasse, A., Ageorges, A., Atanassova, R., Barrieu, F.,
Couture, C., Dedaldechamp, F., Delrot, S., Glissant, D., Grimplet, J.,
Hamdi, S., Romieu, C. and Terrier, N.
Generation of Expressed Sequence Tag from Grape Berry (skin, pulp
or seeds) at Various Developmental Stages
Unpublished (2002)
                                                                                                                                                                  plasmid inserts
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UMR 619 - Equipe Biologie de la Vigne
Universite de Bordeaux I, Institut National de la Recherche
4112, USA
401 585 5606
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71, Avenue
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                                                             Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Vitis vinifera"
/mol_type="mRNA"
/cultivar="Cabernet Sauvignon"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /dev_stage="ripening stage"
/clone_lib="Ripe Grape Skin Triplex2 Library"
/note="Organ: Fruit skin; Vector: Lambda TriplEx2; Site_1:
SfiIA; Site_2: SfiIB; Oriented library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /db_xref="taxon:29760"
/clone="B3CS00RL004E11"
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mid UUGC1M library Mus musculus
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Sciurognathi; Muridae; Murinae; Mus
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AUTHORS
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AZ809971/c
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VERSION
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               Unpublished (2000)
Contact: Robert B. We
University of Utah Ge
University of Utah
Rm. 308, Biomedical I
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                                                                                                                                                                                                                                                                    Mus musculus
Eukaryota; Metazoa; Chordata;
Bukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
1 (bases 1 to 28)
1 (bases 1 to 28)
1 Arber, M.
                                                                                                                                               Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.

Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AZ809971
28 bp UNA
2M0074C14F Mouse 10kb plasmid UUGCIM library Mu
clone UUGC2M0074C14 F, genomic survey sequence.
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Class: plasmid ends
High quality sequence stop: 28.
Location/Qualifiers
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Insert Length: 10000 Std Erry
Plate: 0303 row: L column: 2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pDNA2 (gi |4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
USA
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100.0%; Pred. No. 1.8e+03;
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                 Polymers Research Bldg.,
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AL038989
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Best Local Similarity
Matches 19; Conserv
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DKFZp566B184_rl 566 (synonym:
DKFZp566B184, mRNA sequence.
                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 29)
                                                                                                                                                                                                                                                                                                                                                                                                    AL038989.1 GI:49682230
EST.
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High quality sequence stop: 28.
Location/Qualifiers
                                                                                                                                                                                                EST (Bloecker, et al.)
Unpublished (1999)
                                                                                                                                                                                                                                                              Bloecker, H., Boecher, M.,
                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0074 row: C column: 1-
                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                          Ingolstaedter Landstr.1, D-85764 Neuherberg, Germany.
   Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWH42 (gi 4732114 |gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="DKFZp566B184"
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/strain="C57BL/6J"
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100.0%; Pr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo
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                                                                                                                                  DKFZp566K043 rl 566 (synonym: DKFZp566K043, mRNA sequence. AL048747
1 (bases 1 to 29)
Koehrer, K., Beyer, A., Mewes, H.W.,
EST (Koehrer, et al.)
                                           Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates;
                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DKFZp566F234 rl 566 (synonym: DKFZp566F234, mRNA sequence.
                                                                                                                                                                               AL048747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ingolstaedter Landstr.1, D-85764 Neuherberg, Germany.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: MIPS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Wiemann,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bloecker, H., Boecher, M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AL039052
                                                                                                                    AL048747.1 GI:4727818
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AL039052.1 GI:49682239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AL039052
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                                                                                      Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                            Similarity
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                                                                                                                                                                                                                                                     CAAAAAAAAAAAAAAA 28
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ilarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Homo sapiens"
/mol_type="mRNA"
/mol_type="mRNA"
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/clone="DKFZp566F234"
/tlssue_type="kidney"
/dev_stage="fetal"
/lab_host="Xl-2blue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /tissue_type="kidney"
/dev_stage="fetal"
/lab_host="xl-2blue"
/clone_lib="566 (synonym: hfkd2)"
/note="Vector: pAMP1; Site_1: Not:
                                                                                                                                                                                                                                                                                                                                                                                   'clone_lib="566 (synonym: hfkd2)"
'note="Vector: pAMP1; Site_1: NotI; Site_2:
                                                                                                                                                                                                                                                                                                             0.7%; Score 19; DB 1; L
100.0%; Pred. No. 1.8e+03;
htive 0; Mismatches 0;
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Pred. No.
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: hfkd2) Homo sapiens cDNA clone
                                            Craniata; Ve
Catarrhini;
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1.8e+03;
                Gassenhuber, J. and Wiemann, S.
                                            Vertebrata; Euteleostomi; i; Hominidae; Homo.
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                                                                                                                                                                               EST 04-SEP-2003
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BQ590537/c
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Best Local S
Matches 19
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                                                                                                                                                                                                                                                                                                                                                               Email: weisshaa@mpiz-koeln.mpg.de
Insert Length: 29 Scd Error: 0.00
Plate: 19 row: C column: 03
Seq primer: T7; GTAATACGACTCACTATAGGGC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Max-Planck-Institute for Plant Breeding Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Construction of a 'unigene' cDNA clone set by oligonucleotide fingerprinting allows access to 25 000 potential sugar beet go Plant J. 32 (5), 845-857 (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryotā; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Caryophyllales; Amaranthaceae; Beta.

1 [bases 1 to 29]
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fax: 00492215062851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Herwig, R., Schulz, B., Weisshaar, B., Hennig, S., Steinfath, M., Drungowski, M., Stahl, D., Wruck, W., Menze, A., O'Brien, J., Lei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Beta vulgaris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADIS DNA core facility at MPIZ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Weisshaar B
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Homo sapiens"
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/clone_tib="MPIZ-ADIS-024-storage root"
/note="Vector: pCMVSPORT6; Site_1: SalI; Site_2: NotI;
/note="Vector: pCmVSPORT6; Site_1: SalI; Site_2: NotI;
/cDNA library from sugar beet, library provided by KWS
Kleinwanzlebener Saatzucht AG Binbeck, Germany, contact:
kleinwanzlebener Saatzucht AG Binbeck, Germany, contact:
b.schulz@kws.de; cloning sites SalI-NotI, primer sites and
                                                                                                                                                                                                                                            /organism="Beta vulgaris"
/mol_type="mRNA"
/cultivar="KWS2320 (double haploid, monogerm breeding
                                                                                                                                                            /db_xref="taxon:161934"
/clone="024-019-C03"
                                                                                                                                                                                                          db_xref="GABI:189579"
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Pred. No.
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1.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            000 potential sugar beet genes
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VERSION
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BX551339/c
LOCUS
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Best Local
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                                         19;
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                                                                                                                                                                                                                                                                                                                                                                                               Contact: Hall N
Pathogen Sequencing Unit
The Sanger Institute The Wellcome Trust Genome Campus
Hinxton, Cambridge, CB10 1SA; UK
Request for clones, please contact: Mike Lehane
Prof. M.J.Lehane
School of Biological Sciences,
University of Wales,
Bangor LL57 2UW
All clones with suffix qlc are reverse primer reads st
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Glossina morsitans morsitans
Glossina morsitans morsitans
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        response genes
Genome Biol. 4 (10), R63 (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lehane, M.J., Aksoy, S., Gibson, W., Kerhornou, A., Berriman, M., Hamilton, J., Soares, M.B., Bonaldo, M.F., Lehane, S. and Hall, N. Adult midgut expressed sequence tags from the tsetse fly Glossina morsitans morsitans and expression analysis of putative immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hippoboscoidea; Gl
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BX551339 Glossina morsitans morsitans adult infected gut Glo
morsitans morsitans cDNA clone Tsel18a05_plc, mRNA sequence.
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САЛАЛАЛАЛАЛАЛАЛАЛ 2679
                                                                                                                                                                                                                                                                                                                                                                            clones with suffix qlc are reverse of the cDNA all plc reads are from
                                 0.7%; Sollarity 100.0%; I Conservative 0;
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                                                                                                                                                                                                                                                            /mol_type="mRNA"
/sub_specion"
                                                                                                                                 /note="country: Zimbabwe;
T.brucei"
                                                                                                                                                                                 /sub_species="morsitans"
/db_xref="taxon:37546"
/clone="Tse11805_plc"
/tissue_type="adult infected gut"
/clone_Tib="Glossina morsitans morsitans adult infected
                                                                                                                                                                                                                                                                                     /organism="Glossina morsitans
/mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SP6-Sall-CCACGCGTCCG-Sprime-cDNA-polyA-CC-NotI-T7; Note Sequencing granted in the context of the GABI-Beet project, local PI: Dr. Katharina Schneider, coordinator: Prof. Christian Jung; Sequence submission managed by RZPD/GABI-Primary database: http://gabi.rzpd.de*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          orientation:
                                                                                                                                                                                                                                                                                                                                        ocation/Qualifiers
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b; Pred. No. 1.8
0; Mismatches
                                                       Score 19; 
; Pred. No.
                                     Mismatches
                                                       DB 5; Le
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                                                                      Length 29;
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gut Glossina
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BX555108/c
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Glossina morsitans morsitans
Glossina morsitans morsitans
Glossina morsitans morsitans
Eukaryota; Metazoa; Arthropoda; Hexapoda; Inse
Eukaryota; Endopterygota; Diptera; Brachycera;
Neoptera; Endopterygota; Glossina.
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EX555108 Glossina morsitans morsitans adult infected gut Glossina
morsitans morsitans cDNA clone Tse19e05_plc, mRNA sequence.
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School of Biological Sciences,
University of Wales,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Hall N
Pathogen Sequencing Unit
The Sanger Institute The Wellcome Trust Genome
Hinxton, Cambridge, CB10 1SA, UK
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Lehane,M.J., Aksoy,S., Gibson,W., Kerhornou,A., Berriman,M.,
Hamilton,J., Soares,M.B., Bonaldo,M.F., Lehane,S. and Hall,N.
Adult midgut expressed sequence tags from the tsetse fly Glossina
     Lehane,
                   Hippoboscoidea; Glossinidae; Glossina.
1 (bases 1 to 29)
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygo
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
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Glossina morsitans morsitans
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1 (bases 1 to 29)
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  of the cDNA all plc reads are from
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/tissue_type="adult infected gut"
/clone_lib="Glossina morsitans morsitans adult infected
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|mol_type="mRNA"
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Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Neoptera; Glossinidae; Glossina.
                                                                    Pathogen Sequencing Unit
The Sanger Institute The Wellcome Tr
Hinxton, Cambridge, CB10 1SA, UK
Request for clones, please contact:
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                                                                                                                                                                                                      response genes
Genome Biol. 4 (10),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BX555418 Glossina morsitans morsitans adult infected morsitans morsitans cDNA clone Tse20d06_plc, mRNA sec
                 School of Biological Sciences,
University of Wales,
                                                         Prof. M.J.Lehane
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29 bp

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linear

sequence.

gut Glossina

R63

(2003)

Kerhornou, A.,

Berriman, M.

Pterygota;

Wellcome Trust Genome

Mike Lehane

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Prof. M.J.Lehane
School of Biological Sciences,
University of Wales,
Bangor LL57 2UW
2661 САЛАЛАЛАЛАЛАЛАЛА 2679
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Pathogen Sequencing Unit
The Sanger Institute The Wellcome Trust Genome Campus
Hinxton, Cambridge, CB10 1SA, UK
Request for Clones, please contact: Mike Lehane
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/tissue_type="adult infected gut"
/clone_Tib="Glossina morsitans morsitans
                                                                                                                                note="country: Zimbabwe;
|.brucei"
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db_xref="taxon:37546"
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Sequence 30 from Patent W003072818.
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                                  0.7%; Score 20; DB
larity 100.0%; Pred. No. 44,
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/mol type="unassigned DNA"
/mol type="taxon:32630"
/mbce="Beschreibung der kuenstl
/noce="Beschreibung der kuenstl
                                                                                              note="LNA-T (Locked Nucleic Acid)"
mod_base=OTHER
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AX825158/c
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AX825134/c
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2660 ACAAAAAAAAAAAAAAAAA 2679
            Boekenkamp, D., Dieck, T.H. and Hoppe, H.U. Method for sorting single-stranded nucle Patent: WO 03072818-A 56 04-SEP-2003; Degussa Bioactives GmbH (DE)
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larity 100.0%; Pred. No.
Conservative 0; Mismatc
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/note="Beschreibung der kuenstlichen
Sequenz:Capture-Oligonukleotid"
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Location/Qualifiers
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RESULT 7
AR051244
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                             Sequence 12
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Pickup, D.J., Patel, D. and Antczak, J.B.
Site-specific RNA cleavage
Patent: US 5578468-A 42 26 NOV-1996;
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                            Unclassified.
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/note="Beschreibung der kuenstlichen
Sequenz:Capture-Oligonukleotid"
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/mod_base=OTHER
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/mod_base=OTHER
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100.0%; Pred. No. 44
tive 0; Mismatches
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                                                                              Convergent synthesis of branched and multiply connected macromomolecular structures
Patent: US 5571677-A 12 05-NOV-1996;
                                                                                                                                                                                        Sequence 12 from patent US I28373 I28373.1 GI:1819149
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                                                                                                                    Gryaznov,S.M.
                                                                                                                                                       Unknown.
                                                                                                                                                                   Unknown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unclassified.
1 (bases 1 to 30)
Gryaznov,S.M.
Convergent synthesis of branched and multiply connected macromolecular structures
Patent: US 5830658-A 12 03-NOV-1998;
                                                                                                                                            Unclassified.
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Sequence 12 from patent
AR127791
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Similarity 100.0%; Pred. No.
20; Conservative 0; Mismatch
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ilarity 100.0%;
Conservative (
                                                                      Location/Qualifiers
                                 /organism="unknown"
/mol_type="unassigned
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/mol_type="unassigned
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AR140557/c
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AR140279/c
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Sequence 32
AR139960
AR140557 20 bp
Sequence 32 from patent US 6207802.
AR140557.1 GI:14483053
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Zsebo, K.M., Bosselman, R.A., Suggs, S.V. and Martin, F.H.

Method for enhancing the effciency of gene transfer with stem
factor (SCF) polypeptide
Patent: US 6207454-A 32 27-MAR-2001;
Location/Qualifiers
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                                                                                                                                             l Similarity
19; Conserva
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Zsebo, K.M., Bosselman, R.A., Suggs, S.V.

DNA encoding stem cell factor

Patent: US 6207417-A 32 27-MAR-2001;

Location/Qualifiers
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llarity 100.0%;
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/mol_type="unassigned DNA"
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/mol_type="unassigned
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Sequence 17 from Patent WO03072818.
AX825119
                                    . Similarity
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Zsebo, K.M., Bosselman, R.A., Suggs, S.V.
Stem cell factor and compositions
Patent: US 6207802-A 32 27-MAR-2001;
Location/Qualifiers
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larity 100.0%; Pred. No.
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                                                                                                                  /note="LNA-T (Locked
/mod_base=OTHER
18
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Sequenz:Capture-Oligonukleotid"
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/db xref="taxon:32630"
/note="Taxon-"
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                                                                                         /note="LNA-T (Locked
/mod_base=OTHER
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/mod_base=OTHER
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/mol_type="unassigned
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AX825120/c
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                                      Boekenkamp, D., Dieck, T.H. and Hoppe, H.U. Method for sorting single-stranded nucle Patent: WO 0307/2818-A 19 04-SEP-2003; Degussa Bioactives GmbH (DE)
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Sequence 19 from Patent WO03072818.
AX825121
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AX825120
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(larity 100.0%; Pred. No. 1.7e+02;
Conservative 0; Mismatches 0;
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/mol type="unassigned DNA"
/db xref="reaxon:32630"
/noTe="Beschreibung der kuenstlichen
sequenz:Capture-Oligonukleotid"
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/mod_base=OTHER
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Sequence 20 from Patent WO03072818.
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/mol type="unassigned DNA"
/db_xref="reaxon:32630"
/note="Beschreibung der kuenstlichen
Sequenz:Capture-Oligonukleotid"
                            note="LNA-T (Locked Nucleic Acid)"
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Sequence 21 from Patent WO03072818.
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 AX825124
Sequence 22 from
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larity 100.0%; Pred. No. 1.7e+0
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/note="Beschreibung der kuenstlichen
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                                                                                                                             Boekenkamp,D., Dieck,T.H. and Hoppe,H.U. Method for sorting single-stranded nucleic acids Patent: WO 03072818-A 23 04-SEP-2003; Degussa Bioactives GmbH (DE)
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/bound_moiety="Biotin"
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/note="Beschreibung der kuenstlichen
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/mol_type="unassigned DNA"
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larity 100.0%; Pred. No. 1.7e+02;
Conservative 0; Mismatches 0;
         /note="LNA-T (Locked Nucleic Acid)"
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                                        /note="LNA-T (Locked
/mod_base=OTHER
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mod_base=OTHER
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mod_base=OTHER
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from Patent W003072818.
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AX825128/c
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Best Local Similarity 100.0%; Pred. No. 1.7e+02;
Matches 19; Conservative 0; Mismatches 0;
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Sequence 26 from Patent WO03072818.
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1 Similarity 100.0%; I
19; Conservative 0;
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sequenz:Capture-Oligonukleotid"
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25 from Patent WC03072818.
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Pred. No. 1.7e+02;
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Matches 19; Conserv
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/mol type="unassigned DNA"

/db_xref="taxon:32630"

/note="Beschreibung der kuenstlichen

sequenz:Capture-Oligonukleotid"
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Sequenz:Capture-Oligonukleotid"
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Sequence 28 from Patent WO03072818.
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sequenz:Capture-Oligonukleotid"
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100.0%; Pred. No. 1.
tive 0; Mismatches
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AX825156/c
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                                                       Bookenkamp, D., Dieck, T.H. and Hoppe, H.U. Method for sorting single-stranded nucle Patent: WO 03072818-A 54 04-SEP-2003; Degussa Bioactives GmbH (DE)
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AX825156
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Sequence 53 from Patent
AX825155
AX825155.1 GI:39750884
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ilarity 100.0%; Pred. No. 1.7e+02;
Conservative 0; Mismatches 0;
         organism="synthetic construct"
                                       Location/Qualifiers
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/mod_base=OTHER
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Sequenz:Capture-Oligonukleotid"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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AX825157/c
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Location/Qualifiers
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/note="Beschreibung der kuenstlichen
Sequenz:Capture-Oligonukleotid"
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/mod_base=OTHER
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/note="LNA-T (Locked Nucleic Acid)"
                            /note="LNA-T (Locked Nucleic Acid)"
/mod_base=OTHER
                                                                       /note="LNA-T (Locked Nucleic Acid)"
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/mod_base=OTHER
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Sequenz:Capture-Oligonukleotid"
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/db_xref="taxon:32630"
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Sequence 62 from Patent
AX825164
   BD085544
Method of comparison
BD085544
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Sequenz:Capture-Oligonukleotid"
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/mod_base=OTHER
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db_xref="taxon:32630"
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UNITECH CO LTD
OS Homo sapiens (human)
PN JP 2001333800-A/1
PD 04-DEC-2001
PF 30-MAY-2000 JP 2000160324
PI KAORI SHIMADA
PC C1201/68,C12N15/09,G01N33/50,C12N15/00
CC Method of comparison and detection of RNA amounts
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synthetic construct
artificial sequences.
1 (bases 1 to 23)
1 Hartwich,G. and Heller,A.
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OS Artificial Sec
PN JP 2002532386-
PD 02-OCT-2002
PF 19-NOV-1999 JI
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Method of
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G01N27/416,G01N27/48,G01N33/483,G01N33/50,G01N33/566,C12N15/00,
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                                                                                                                                                                                         PF 19-NOV-1999 JF 2000583928
PR 23-NOV-1998 DE 198 53 957.6,29-APR-1999 DE 199 21 940.0 PI
GERHARD HARTWICH, ADAM HELLER
PC C07H21/00,C07H21/02,C07H21/04,C12N15/09,C12Q1/68,G01N27/12,
                                                                                                                                                                                                                                                                                                                       Method of electrochemically detecting nucleic Patent: JP 2002532386-A 16 02-OCT-2002;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                         Method of electrochemically detecting nucleic acid Location/Qualifiers
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JP 2002532386-A/16
02-OCT-2002
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C12Q1/68,C12N15/09,G01N33/50,C12N15/00
Method of comparison and detection of RNA amount and DNA
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/db_xref="taxon:32630"
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/mol_type="genomic RNA"
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detecting nucleic acid
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17-JUL-2003

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and DNA amount

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RESULT 32
AR431312/c
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BD196419/c
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PC C12N1/21,C12N5/10,C1
,C12N15/00,C12N15/00,
PC C12N5/00,C12N15/00
CC primer oligonuclecti
FH Key
FT misc_binding 1.
                                           Unclassified.

1 (bases 1 to 24)

Vaisberg,E.A., Adams,C.L., Sabry,J.H. and Crompton,A.M.

Database system including computer code for predictive cellular
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19; Conserv
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OS Homo sapiens (human)
PN JP 2002516657-A/8
PD 11-JUN-2002
PP 22-DEC-1998 JP 2000525562
PF 22-DEC-1997 US 08/996306,09-SEP-1998 US 60/099658 PI
DANIEL COHEN, MATRA BLUMENFELD, ILYA CHUMAKOV, LYDIE BOUGUELERET PC
C12N15/09,C12N15/09,A01K67/027,C07K14/47,C07K16/18,C12N1/15, PC
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                     Patent: US 6651008-A 6 18-NOV-2003;
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AR431312
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                                                                                                                     Unknown
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JP 2002516657-A/8.
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larity 100.0%; Pred. No. 1.7e+02;
Conservative 0; Mismatches 0;
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Location/Qualifiers
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            Location/Qualifiers
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                                                                                                                                              GI:40193280
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AX817782
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              misc_feature
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Sequence 8 from Patent WO02068627
AX838369
                                                                                                                                                                                   synthetic construct synthetic construct
                                                                                                                 Patent:
                                                                                                                                                                   artificial sequences.
                                                                                                                             Vector constructs
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synthetic construct
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                                                                                                             WO 02068627-A 8 06-SEP-2002;
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/mol_type="genomic [
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                                       organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
'note="Fig. 1C. SV40 early Poly(A) site"
                             'note="Viral vector
                                                                                                  location/Qualifiers
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O 02067861-A 18 06-SEP-2002;
Location/Qualifiers
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/db_xref="taxon:32630"
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BD161931/c
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AUTHORS
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BD097127/c
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artificial sequences.

Et 1 (bases 1 to 24)

Et 1 (bases 1 to 24)

RS Tanga, M., Okamura, H., Takagi, K. and Takahashi, K.

RS Tanga, M., Okamura, H., Takagi, K. and Takahashi, K.

RS Tanga, M., Okamura, H., Takagi, K. and Takahashi, K.

RS Tanga, M., Okamura, H., Takagi, K. and Takahashi, K.

RS Tanga, M., Okamura, H., Takagi, K. and Takahashi, K.

ROYO KOHAN CO LTD, MICHIFUMI TANGA, HIROSHI OKAMURA, KENICHI TAKAGI,

KOJIRO TAKAHASHI

OS ARTIFICIAL Sequence

PN WO 0155365-A/1

PD 02-AUG-2001 WO 2001JP000443

PF 24-JAN-2001 WO 2001JP000443

PR 27-JAN-2000 JP 00P 019301

PR 27-JAN
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artificial sequences.

1 (bases 1 to 24)
Tanga,M., Okamura,H. and Takahashi,K.
Method for carrying out thermal cycle of
Patent: JP 2002191369-A 8 09-JUL-2002;
TOYO KOHAN CO LTD,KOJIRO TAKAHASHI
OS Artificial Sequence
                                                                                                                                                                                                     JP 2002191369-A/8.
                                                                                                                                                                                                                      BD161931
BD161931.1 GI:27867689
                                                                                                                                                                                                                                                                                   Method for
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WO 0155365-A/1.
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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carrying out thermal
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100.0%; Pred. No.
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of PCR using
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                                                                               PCR
                                                                           using
                                                                               DNA-immobilized
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DNA-immobilized
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AX394507
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SOURCE
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2661 САЛАЛАЛАЛАЛАЛАЛАЛ 2679
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A method for generating hypermutable organisms Patent: WO 0188192-A 4 22-NOV-2001;
The Johns Hopkins University School of Medicine Inc. (US); Nicolaides, Nicholas, C. (US); Sass Grasso, Luigi (US); Vogelstein, Bert (US)
Location/Qualifiers
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Sequence 4 from Patent WO0188192.
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PD 09-JUL-2002
PF 27-DEC-2000 JP 2000399573
PI MICHIFUMI TANGA, HIROSHI OKAMURA, KOJIRO TAKAHASHI
PI MICHIFUMI TANGA, C12Q1/68, C12X15/00, C12X15/00 CC
C12X15/09, C12X15/09, C12Q1/68, C12X15/00, C12X15/00 CC
Carrying out thermal cycle of PCR using DNA- CC
immobilized
                                                                Sequence 52 from Patent AX394507
synthetic
               synthetic construct
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Location/Qualifiers
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/mal_tyne="unassigned DNA"
/mal_tyne="taxon:32630"
/mb[="Recombinant DNA"
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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3, Philip, M. (US)
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BD192375/c
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Manmalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 26)

Medel, P. A.B., Cohen, M., Colpitts, T.L., Friedman, P.N., Gordon, J., Granados, E.N., Hodges, S.C., Klass, M.R., Kratochvil, J.D., Russell, J.C., Scheffel, C.P., Stroupe, S.D. and Yu, H. Resgents and methods useful for detecting diseases of the breast Patent: JP 2002516576-A 14 04-JUN-2002; ABBOTT LABORATORIES PN JP 2002516576-A/14

PD 04-JUN-2002
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Reagents and methods useful f
BD192375
BD192375.1 GI:33002114
JP 2002516576-A/14.
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Sequence 59 from Patent WO0218638.
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/db_xref="taxon:32630"
/note="Synthetic oligonucleotide"
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/mol_type="unassigned DNA"
/db xref="taxon.32630"
/noTe="Synthetic oligonucleotide"
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Sequence 3
AX338547
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PI STROUPE
PI HONG YU
PC C12N15/
G01N33/574
CC Strande
CC Topolog
FH Key
Nicolaides, N.C., Kinzler, K.W.
                                                       synthetic construct synthetic construct
                                         artificial sequences.
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Patent: US 5707807-A 3 13-JAN-1998;
Location/Qualifiers
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C12N15/12,C07K14/47,C12Q1/68,C12N15/85,C12N5/10,C07K16/18,
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20-JUN-1997 US 08/879354
PATRICIA A BILLING MEDEL, MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N FRIEDMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Topology: Linear;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              JULIAN GORDON, EDWARD N GRANADOS, STEVEN C HODGES, MICHAEL R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     JON D KRATOCHVIL, JOHN C RUSSELL, CHRISTI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                    /organism="unknown"
/mol_type="unassigned
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Mus sp."
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                                                                                                                        26 bp
from Patent WO0188192.
                                                                                                GI:18128947
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100.0%; Pred. No.
tive 0; Mismatc
                                                                                                                                                                                                                                                                               0.7%; Score 19; DB 6;
100.0%; Pred. No. 1.7e+02
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           Sass, P.M.,
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           Grasso,
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thes 0;
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          Vogelstein,
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RESULT 44
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synthetic construct
artificial sequences.

El (bases 1 to 27)

El (bases 1 to 27)

RS Tanga, M., Okamura, H., Takagi, K. and Takahashi, K.

RS Tanga, M., Okamura, H., Takagi, K. and Takahashi, K.

Support for immobilizing nuclectide and process for producing the Patent: WO 0153365-A, 2 02-AUG-2001,

KOJIRO TAKAHASHI

OS ARTIFICIAL Sequence
PN WO 015535-A/2

PP 02-AUG-2001 WO 2001JP000443

PF 24-JAN-2001 WO 2001JP000443

PF 24-JAN-2000 JP 00P 019301

PF 24-JAN-2000 JP 00P 019301

PG C12N15/10,C07H21/04//G01N33/50,C1201/68

CC Support for immobilizing nuclectide and process for producing the same

CC Support for immobilizing nuclectide and process for producing the same

FH Key

I. Castion/Onslifices

FH Key

I. Castion/Onslifices

FI Source

//organism='Artificial Sequence'.
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BD161932
Method for
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WO 0155365-A/2.
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Patent: WO 0188192-A 3 22-NOV-2001;
The Johns Hopkins UNiversity
Inc. (US); Nicolaides, Nicholas, C. (US); Sass, Philip, M. (US);
Grasso, Luigi (US); Vogelstein, Bert (US)
Location/Qualifiers
                                                                                                                                                             Similarity
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                                                                                                                                                                                                                                                                  /organism='Artificial
                                                                                                                                                                                                                 /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note="Recombinant DNA"
            carrying out
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       immobilizing
                                                                                                                                                           0.7%; Score 19;
100.0%; Pred. No.
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100.0%; Pred. No. 1.
ive 0; Mismatches
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           27 bp
thermal
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           linear
PCR using
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            PAT 17-JAN-2003
DNA-immobilized
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                          2661 САЛАЛАЛАЛАЛАЛАЛА 2679
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent: JY ZVOLLOW TOYO KOHAN CO LTD, KOJIRO TAKAHASHI
OS Artificial Sequence
PN JP 2002191369-A/9
PD 09-JUL-2002
PF 27-DEC-2000 JP 2000399573
PI MICHIFUMI TANGA, HIROSHI OKAMURA, KOJIRO TAKAHASHI
C12N15/09,C12N15/09,C12Q1/68,C12N15/00,C12N15/00 CC
5
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Sequence 10 from Patent
AX391845
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  S
                                                         Similarity 100. 19; Conservative
                                                                                                                                                                                                                       Reimholz,R. and Ploeger,F. Method for identifying peptides that can and the use of peptide sequences of this Patent: WO 0216574-A 10 28-FEB-2002, Xzillion GmbH & CO.KG (DE)
                                                                                                                                                                                                                                                                                                                                synthetic construct
synthetic construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 (bases 1 to 27)
Tanga, M., Okamura, H. and Takahashi, K.
Method for carrying out thermal cycle
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synthetic construct
artificial sequences.
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UP 2002191369-A/9.
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                                                                                                                                /organism="synthetic construct"
/mal_type="unassigned RNA"
/db xrefs"ttaxon:32630"
/noEe="Puromycin-Linker-RNA-Teil"
                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="synthetic co
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/db_xref="taxon:32630"
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                                                     0.7%; Score 19; DB 100.0%; Pred. No. 1. tive 0; Mismatches
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100.0%; Pred. No. 1.7
tive 0; Mismatches
28
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hes 0;
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REFERENCE
AUTHORS
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BD165919
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AX430216
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AR438517
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                                                 BD165919

Method for melting curve a
BD165919

BD165919.1 GI:27871731

JP 2002191384-A/7.

unidentified
     unclassified.
1 (bases 1 to :
Dietmaier, W.
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19; Conserv
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                                         unidentified
                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                        Method for melting curve analysis of repetitive pcr products Patent: EP 1207210-A 7 22-MAY-2002; Roche Diagnostics GmbH (DE); P. HOPFMANN-LA ROCHE AG (CH)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29 bp
Sequence 7 from Patent EP1207210.
AX430216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AX430216.1 GI:21655581
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1 (bases 1 to 29)
Dietmaier, W.
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Sequence 7 from
AR438517
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.7%; Score 19; DB 6; Lilarity 100.0%; Pred. No. 1.8e+02; Conservative 0; Mismatches 0;
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/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                 ocation/Qualifiers
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/mol_type="genomic DNA"
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Pred. No. 1.8e+02;
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AX079108
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Sequence 7 from Patent WO0106226.
AX079109
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Sequence 6 from Patent WO0106226.
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AX079108.1 GI:13158682
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/note="Oligonukleotid"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="unidentified"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism='Homo
                                                                                                                                                          0.7%; Sur
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                                               GI:13158683
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AR034896/c
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Process for preparing polynucleotides
tightly packed bed
Patent: US 5869643-A 12 09-FEB-1999;
Location/Qualifiers
                                                                                                                1 (bases 1 to 18)
Chatelain, F. and Kumarev, V.
                                                                                                                                                     Unknown
                                                                              Process for preparing polynucleotides tightly packed bed Patent: US 5869643-A 18 09-FEB-1999;
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/mol_type="unassigned
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/mol type="unassigned DNA"
/db xref="taxon:32630"
/noTe="Oligonukleotid"
                                  /organism="unknown"
/mol_type="unassign:
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AR097579.1
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Sequence 3
AR058305
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Sequence
AR106506
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l Similarity 100.0%; P
18; Conservative 0;
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Lin,C.-I.Patsy., Wallace,R.Bruce., Cossman,J. and French,C.Method and formulation for lyophilizing cultured human cell preserve RNA and DNA contained in cells for use in moleculabiology experiments
Patent: US 6071745-A 9 06-JUN-2000;
Location/Qualifiers
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De Rose, R., Douce, R., Duval, M., Job, C.
Seed specific biotinylated protein, SB
Patent: US 5837820-A 3 17-NOV-1998;
                                                                                                                                                                                                                                                                                                              Unknown
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             30
                                                                                                                                                                                /organism="unknown"
/mol_type="unassigned DNA"
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/mol_type="unassigned
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synthetic construct
artificial sequences.
1 (bases 1 to 18)
Mancharan, M., Cook, P.D., Prakas
Aminooxy-modified nucleoside co
produced therefrom
                                                                                                                 18;
                                                                                                                                                                                                                                                                                                                                                                  Patent: JP 2002522447-A 14 23-JUL-2002;
ISIS PHARMACEUTICALS INC
OS Artificial Sequence
PN JP 2002522447-A/14
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Keeling, P. and Guan, H.
Starch encapsulation
Patent: US 6107060-A 30 22-AUG-2000;
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09-AUG-1999 JP 2000563675
07-AUG-1998 US 09/130973
MUTHIAH MANOHARAN, PHILIP DAN COOK, THAZHA P PRAKASH, ANDREW M
                                                                                                             0.7%;
larity 100.0%;
Conservative (
                                                                                                                                                                                                                                                                Description of Artificial Sequence: antisense sequence FH
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                                                                                                                                                                       /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                                                                                                                                                                                                      /organism='Artificial Sequence'
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100.0%; Pred. No. 6.;
tive 0; Mismatches
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E28536/c
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Kenichi,H., Hiroshi,Y. and Masahide,N.

Method for labeling oligonucleotide and utilization thereof

Patent: JP 1999075880-A 3 23-MAR-1999;

CHEMO SERO THERAPEUT RES INST
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E28536
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Kenichi,H., Hiroshi,Y. and Massahide,N.

Method for labeling oligonucleotide and utilization thereof

Patent: JP 1999075880-A 2 23-MAR-1999;

CHEMO SERO THERAPEUT RES INST
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JP 1999075880-A/2.
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JP 1999075880-A/3
23-MAR-1999
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JP 1999075880-A/2
23-MAR-1999
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                                                                /organism='Unidentified'
Location/Qualifiers
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/mol_type="genomic DNA"
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/db_xref="taxon:32644"
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AR208426/c
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Sequence
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Molecular indexing for expressed gene Molecular indexing for expressed gene Patent: US 5707807-A 16 13-JAN-1998;

Location/Qualifiers
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18; Conserv
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Kaufman, J.C., Roth, M.E., Lizardi, P.M.,
Binary encoded sequence tags
Patent: US 6383754-A 6 07-MAY-2002;
Location/Qualifiers
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/mol_type="unassigned DNA"
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AR412363
AR412363.1
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Kurz,M., Lohse,P. and Wagner,R.

Peptide acceptor ligation methods

Patent: US 6429300-A 24 06-AUG-2002;

Location/Qualifiers
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AR222464
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Lin,C.-I.P., Wallace,R.B., Cossman,J. and French,C.
Method and formulation for lyophilizing cultured human cells
preserve RNA and DNA contained in cells for use in molecular
biology experiments
Patent: US 6410321-A 9 25-JUN-2002;
Location/Qualifiers
                               1 (bases 1 to 18)
Manoharan, M., Cook, P.D., Prakash, T.P
Aminooxy-modified nucleosidic compou
prepared therefrom
Patent: US 6639062-A 14 28-OCT-2003;
                                                                                                  Unknown
                                                                                      Unclassified
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                     Location/Qualifiers
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AR487019
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AR473365/c
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                                                                                                                                       Unclassified.

1 (Dases 1 to 18)
Thirstrup,K., Warthoe,P. and Pettersson,N.B.
Process for amplifying and labeling single stranded ligated adaptor mediated amplification Patent: US 6706476-A 6 16-MAR-2004;
                                                                                                                                                                                                                                                                                                                                                                                  18;
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                                                                                                                                                                                                              Unknown
                                                                                                                                                                                                                                              AR487019.1
                                                                                                                                                                                                                                                         AR487019
Sequence 6 from
AR487019
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   biology experiments
Patent: US 6686460-A 9 03-FEB-2004;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 (bases 1 to 18)
Lin,C. I.P., Wallace,R.B., Cossman,J. and French,C.
Method and formulation for lyophilizing cultured human cells
preserve RNA and DNA contained in cells for use in molecular
                                                                                                                                                                                                                        Unknown.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 9
AR473365
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ilarity 100.0%;
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Conservative 0;
                                                                                                /organism="unknown"
/mol_type="genomic DNA"
                                                                                                                                  Location/Qualifiers
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/mol_type="genomic DNA'
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                                                    Score 18;
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6.3e+02;
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RESULT 68
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AR487020/c
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synthetic construct
synthetic construct
artificial sequence
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1 Similarity 100.0%;
18; Conservative
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Method for isolating anionic organic substances from aqueous
systems using cationic polymer nanoparticles
Patent: WO 9910527-A 4 04-MAR-1999;
SUEDDEUTSCHE KALKSTICKSTOFF (DE); BAYER ERNST (DE)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AX004875 18 bp
Sequence 4 from Patent WO9910527.
AX004875
AX004875.1 GI:9928275
                                                    AX004879.1
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Thirstrup, K., Warthoe, P. and Pettersson, N.B.
Process for amplifying and labeling single stranded cDNA by
ligated adaptor mediated amplification
Patent: US 6706476-A 7 16-MAR-2004;
                                                                                                                                                                                                                                                                                                                                                                                                                                         artificial sequences.
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Similarity 100.0%; I
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                                                                                                                                                                                                                                                                           /organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/note="3' palmityl oligonucleotide"
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/mol_type="genomic DNA"
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from Patent WO9910527
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Pred. No.
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AX008118/c
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AUTHORS
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AX008117
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Sequence 3
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Method for isolating anionic organic substances from
systems using cationic polymer nanoparticles
Patent: WO 9910527-A 8 04-MAR-1999;
SUEDDEUTSCHE KALKSTICKSTOFF (DE); BAYER ERNST (DE)
                                                                                                                                                                   _
                                                                                                                                                                                                                                                                                                                                                     and its analogues
Patent: WO 9967378-A 2 29-DEC-1999;
DAMHA MASSAD JOSE (CA); PARNIAK MICHAEL A (CA); WILDS CHRISTOPHER (CA); UNIV MCGILL (CA); ARION DOMINIQUE (CA); NORONHA ANNE M (CA);
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 2
AX008117
                                   synthetic construct synthetic construct
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  Damha, M.J.,
                                                                       AX008118.1
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/mod_base=um
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="unassigned RNA"
/db_xref="taxon:32630"
/note="2' methyl-modified oligonucleotide"
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from Patent WO9967378
                                                                       GI:9995743
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 Parniak, M.A.,
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100.0%; Pred. No. 6.3e+02;
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WO9967378
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 Wilds, C.,
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AX008123
AX008123.1
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 Borkow, G.
            Damha,M.J.,
                                  artificial
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2662 AAAAAAAAAAAAAAAAA 2679
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Sequence 7 from Patent WO9967378.
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Patent: WO 9967378-A 3 29-DEC-1999;
DAMHA MASSAD JOSE (CA); PARNIAK MICHAEL A (CA); UNIV MCGILL (CA); ARION DOMINIQUE (CA) BORKOW GADI (IL)
                                                                                                                                                                                                                                                                                                                                                                                                                                               and its analogues
Patent: WO 9967378-A 7 29-DEC-1999;
DAWHA MASSAD JOSE (CA); PARNIAK MICHAEL
(CA); UNIV MCGILL (CA); ARION DOMINIQUE
BORKOW GADI (IL)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         artificial sequences.
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                                           construct
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/mol_type="unassigned RNA"
/db_xref="taxon:32630"
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/mol_type="unassigned I
/db_xref="taxon:32630"
/note="Use as an oligon
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from Patent WO9967378.
                              sequences.
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Parniak, M.A.,
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100.0%; Pred. No. 6.:
tive 0; Mismatches
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100.0%; Pred. No.
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Wilds, C.,
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Noronha, A.M.
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Sequence 29
AX028845
Muehlegger, K., Angerer, B., Seela, F., Ankenb
Gumbiowski, K. and Zulauf, M.
High density labeling of dna with modified
nucleotides and dna polymerases used
Patent: WO 0068422-A 21 16-NOV-2000;
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AX047271
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Brugliera, F., Holton, T.A. and Michael, M. Genetic sequences encoding flavonoid pat
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Patent: WO 9967378-A 8 29-DEC-1999;
DAMHA MASSAD JOSE (CA); PARNIAK MICHAEL A (CA); WILDS CHRISTOPHER (CA); UNIV MCGILL (CA); ARION DOMINIQUE (CA); NORONHA ANNE M (CA);
BORKOW GADI (IL)
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Llarity 100.0%; Pred. No. 6.3e+02;
Conservative 0; Mismatches 0;
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db_xref="taxon:32630"
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/mol_type="unassigned DNA"
/db_xrcf="taxon:32630"
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Sequence 23
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Gumbiowski, K. and Zulauf, M.
High density labeling of dna with modified
nucleotides and dna polymerases used
Patent: WO 0068422-A 23 16-NOV-2000,
Roche Diagnostics GmbH (DB)
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                                                                                                                /organism="synthetic construct"
/mol_type="unassigned DNA"
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/note="second fragment of SEQ ID NO:
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/mol_type="unassigned DNA"
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(without alignments)
10417.833 Million cell updates
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Sequence 325, App
Sequence 32, Appl
Sequence 32, Appl
Sequence 433, App
Sequence 654, Appl
Sequence 654, Appl
Sequence 10, Appl
Sequence 10, Appl
Sequence 10, Appl
Sequence 11, Appl
Sequence 11, Appl
                                                                                                                                               Description
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US-09-90-803-646-3
US-09-90-803-647-3
US-09-90-803-647-9-913
US-09-90-90-911-6
US-09-976-479-913
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US-09-976-479-913
US-10-125-295-9
US-10-125-297-1
US-10-123-597-1
 Sequence 4, Appli
Sequence 52, Appli
Sequence 53, Appli
Sequence 14, Appli
Sequence 34, Appli
Sequence 84, Appli
Sequence 913, App
Sequence 914, Appli
Sequence 15, Appli
Sequence 16, Appli
Sequence 17, Appli
Sequence 913, App
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Sequence 16, Appli
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Sequence 919, Appli
Sequence 910, Appli
Sequence 911, Appli
Sequence 911, Appli
Sequence 913, Appli
Sequence 914, Appli
Sequence 915, Appli
Sequence 916, Appli
Sequence 917, Appli
Sequence 918, Appli
Sequence 919, Appli
Seq
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ALIGNMENTS

RESULT 1 US-10-205-309-325 ; Sequence 325, Application US/10205309

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SQUAL 3
-09-005-243-32/c
Sequence 32, Application US/09005243
Patent No. US2002018763A1
GENERAL INFORMATION:
APPLICANT: Zeebo, Krisztina M.
APPLICANT: Bosselman, Robert A.
APPLICANT: Suggs, Sidney V.
APPLICANT: Sugra, Sidney V.
APPLICANT: Martin, Francis H.
TITLE OF INVENTION: Stem Cell Factor
NUMBER OF SEQUENCES: 104
CORRESPONDENCE ADDRESS:
ADDRESSEE: 6300 Sears Tower, 233 South Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LT 2
0-205-309-650/c
0-205-309-650/c
iquence 650, Application US/10205309
iblication No. US20030190635A1
INFORMATION:
REPLICANT: Ribozyme Pharmaceuticals, Inc.
INPLICANT: MCSwiggen, James
INPLICANT: MCSwiggen, Tritarfarance Mediat.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ITLE OF INVENTION: RNA Interference Mediated Inhibition of Alzheimer's Disease Using TITLE OF INVENTION: Interfering RNA
ILE REFERENCE: 900/03
ITRENT APPLICATION NUMBER: US/10/205,309
IRRENT FILING DATE: 2002-10-25
WHER OF SEQ ID NOS: 674
FTWARE: PatentIn version 3.0
ID NO 650
INGTH: 19
FTER RNA
CANATON.
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RRENT FILING DATE: 2002-10-25
RBER OF SEQ ID NOS: 674
TWARE: Patentin version 3.0
ID NO 325
IMGTH: 19
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PLICANT: RIDOZYME Pharmaceuticals, Inc.
PLICANT: RIDOZYME Pharmaceuticals, Inc.
PLICANT: McSwiggen, James
PLICANT: McSwiggen, James
PLICANT: McSwiggen, James
PLIE OF INVENTION: RNA Interference Mediated Inhibition of Alzheimer's Disease Using RDF INVENTION: Interfering RNA
E REFERENCE: 900/033
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GANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HER INFORMATION: Description of Artificial Sequence: 205-309-650
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100.0%; Pred. No. 1.7e+02;
ive 0; Mismatches 0;
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1.7e+02;
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MOLECULE TYPE:
US-09-005-243-32
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Best Local Similarity
Matches 19; Conserv
                                                                                                                                  Sequence 32, Application US/09224683
Patent No. US20020031491A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: 25-3856
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                           APPLICANT: Zsebo, Krisztina M. APPLICANT: Bosselman, Robert A. APPLICANT: Suggs, Sidney V. APPLICANT: Martin, Francis H. TITLE OF INVENTION: Stem Cell F. NUMBER OF SEQUENCES: 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 16-OCT-1989
ATTORNEY/AGENT INFORMATION:
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FILING DATE: 01-OCT-1990
PRIOR APPLICATION DATA:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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                             CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 01
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FILING DATE: 24-AUG-1990
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               ADDRESSEE:
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100.0%; Pr
                                             Stem Cell Factor: Composition Claims: 104
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Tower, 233
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; Pred. No.
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   Gerstein, Murray & Borun
South Wacker Drive
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RESULT 5
US-10-671-395-433/c
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Best Local S
Matches 19
                                                                                                                  Sequence 433, Applic Publication No. US20 GENERAL INFORMATION
                                              APPLICANT: Gierse, James K
TITLE OF INVENTION: ANTISENSE MODULATION OF MICROSOMAL PROSTAGLANDIN E2 SYNTHASE
TITLE OF INVENTION: EXPRESSION
CURRENT APPLICATION NUMBER: US/10/671,395
CURRENT FILING DATE: 2003-09-25
                                                                                    APPLICANT: Pharmacia Corp
APPLICANT: Gierse, James
                                  FILE REFERENCE: 1179/1/US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE:
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ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: 07/589,701
FILING DATE: 01-OCT-1990
PRIOR APPLICATION DATA:
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TELEFAX: 25-3856
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nes 19; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 07/9
FILING DATE: 25-NOV-1992
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                           2661 СААЛАЛАЛАЛАЛАЛАЛА 2679
                                                                                                                                    433, Application US/10671395 ion No. US20040132063A1
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                                                                                                                                                                                                                                                                                                                                               DB 9;
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CURRENT APPLICATION NUMBER: US/10/671,395
CURRENT FILING DATE: 2003-09-25
PRIOR APPLICATION NUMBER: 60/413,549
PRIOR FILING DATE: 2002-09-25
NUMBER OF SEQ ID NOS: 1809
SOFTWARE: PatentIn version 3.2
SEQ ID NO 654
LENGTH: 20
TYPE: DNA
                                                                                                                                                                                                                                                   US-10-175-608-32/c
; Sequence 32, Application US/10175608
; Publication No. US20040181044A1
; GENERAL INFORMATION:
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J Sequence 654, Application US/10671395

Publication No. US20040132063A1

GENERAL INFORMATION:

APPLICANT: Gierse, James K

TITLE OF INVENTION: ANTISENSE MODULATION OF MICROSOMAL PROSTAGLANDIN E2 SYNTHASE

TITLE OF INVENTION: EXPRESSION

TITLE OF INVENTION: EXPRESSION

TITLE OF INVENTION: EXPRESSION

TITLE OF INVENTION: EXPRESSION
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PRIOR FILING DATE: 2002-09-25
NUMBER OF SEQ ID NOS: 1809
SOFTWARE: PATENTIN VERSION 3.2
SEQ ID NO 433
LENGTH: 20
TYPE: DIA
ORGANISM: artificial
PEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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Best Local Similarity
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OTHER INFORMATION: Human PGE2 antisense
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                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Marshall, O'Tou
                                                                                                                                              APPLICANT: Zsebo, Krisztina M.
Bosselman, Robert A.
Suggs, Sidney V.
Martin, Francis H.
TITLE OF INVENTION: Stem Cell Factor
NUMBER OF SEQUENCES: 104
COMPUTER READABLE FORM
                                                                                                                                                                                                                                    APPLICANT:
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                                                       STATE: Illinois
               COUNTRY: United States of ZIP: 60606-6402
                                                                        CITY: Chicago
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100.0%; Pred. No. 1.7
tive 0; Mismatches
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100.0%; Pred. No. 1.7
tive 0; Mismatches
                                                                                    O'Toole, Gerstein, Murray & Borun
ower, 233 South Wacker Drive
                                    America
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hes 0;
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RESULT 8
US-10-410-031-190/c
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                                                SOFTWARE: PatentIn version 3.2
SEQ ID NO 190
LENGTH: 21
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Best Local Similarity
Matches 19; Conserv
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APPLICANT: Shockey, Jay M.
APPLICANT: Schnurr, Judy
APPLICANT: Browse, John A
                                                                                                                                                                                                                                                                                                            Sequence 190, Application US/10410031 Publication No. US20040010817A1
                                                                                                              CURRENT APPLICATION NUMBER: US/10/410,031
CURRENT FILING DATE: 2003-04-09
NUMBER OF SEQ ID NOS: 191
                                                                                                                                                                            APPLICANT: Schnurr, Judy
APPLICANT: Browse, John A.
TITLE OF INVENTION: Plant Acyl-CoA Synthetases
FILE REFERENCE: DOW-07654
ORGANISM: Artificial Sequence
                             TYPE: DNA
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INFORMATION FOR SEQ ID NO: 32:
SEQUENCE CHARACTERISTICS:
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COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION UNUBER: US/10/175,608
FILING DATE: 16-Oct-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
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FILING DATE: 07-AUG-2000
APPLICATION NUMBER: 09/486,546
FILING DATE: 24-MAY-1995
APPLICATION NUMBER: 08/172,329
FILING DATE: 21-DEC-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Clough, David W. REGISTRATION NUMBER: 36, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 11-JUN-1990
APPLICATION NUMBER: 07/422,383
FILING DATE: 16-OCT-1989
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APPLICATION NUMBER: 07/537,198
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APPLICATION NUMBER: 09/589,701
FILING DATE: 10-OCT-1991
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FILING DATE: 25-NOV-1992
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BER: 01017/35199
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US-09-853-526-10/c
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; Sequence 10, Application US/09901484A
; Patent No. US20020119460A1
APPLICANT: Bougueleret, Lydie
TITLE OF INVENTION: PROSTATE CANCER GENE
FILE REFERENCE: GENSET.18CPLCP
CURRENT APPLICATION NUMBER: US/09/853,526
CURRENT FILING DATE: 2001-05-11
PRIOR APPLICATION NUMBER: 09/338,907
PRIOR FILING DATE: 1999-06-23
PRIOR APPLICATION NUMBER: 08/996,306
PRIOR APPLICATION NUMBER: 08/996,306
PRIOR FILING DATE: 1997-12-22
                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Cohen, Daniel
APPLICANT: Blumenifeld, Marta
APPLICANT: Ilya, Chumakov
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Best Local :
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SEQ ID NO 10
LENGTH: 24
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PRIOR FILING DATE: 1998-09-09
PRIOR APPLICATION NUMBER: US 09/218,207
PRIOR FILING DATE: 1998-12-22
PRIOR APPLICATION NUMBER: US 09/338,907
PRIOR FILING DATE: 1999-06-23
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CURRENT FILING DATE: 2001-07-09
PRIOR APPLICATION NUMBER: US 08/996,306
PRIOR FILING DATE: 1997-12-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc feature
LOCATION: (1)...(24)
OTHER INFORMATION: primer oligonucleotide PGRT32
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TITLE OF INVENTION: Prostate Cancer Gene
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les 19; Conservative (
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100.0%; Pr/
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; Pred. No.
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; Pred. No.
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                                                                                                                                                                                                               SEQ ID NO 18
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                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/270,922
PRIOR FILING DATE: 2001-02-23
PRIOR APPLICATION NUMBER: US 60/295,037
PRIOR FILING DATE: 2001-66-01
PRIOR APPLICATION NUMBER: US 60/348,670
PRIOR FILING DATE: 2001-66-01
                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE: 4-31704A/GTI
CURRENT APPLICATION NUMBER: US/10/081,969
CURRENT FILING DATE: 2002-02-22
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PRIOR FILING_DATE: 1998-09-09
PRIOR APPLICATION NUMBER: 09/218,207
PRIOR FILING DATE: 1998-12-22
NUMBER OF SEQ ID NOS: 578
                                                                                                                                                                                                                                              PRIOR FILING DATE: 2000-01-14
NUMBER OF SEQ ID NOS: 98
                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Steward, David
APPLICANT: Xie, Yuefeng
TITLE OF INVENTION: No. US20030104625Alel Oncolytic Adenoviral Vectors
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                                                                                                                                                                                                                                SOFTWARE: PatentIn version 3.1
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NAME/KEY: misc_feature
LOCATION: 1..24
OTHER INFORMATION: primer oligonucleotide PGRT32
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TYPE: DNA
ORGANISM: Homo Sapiens
                                                                                                                                                        LENGTH: 24
TYPE: DNA
ORGANISM: Artificial Sequence
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OTHER INFORMATION:
               NAME/KEY: polyA site LOCATION: (3)..(24)
                                                               OTHER INFORMATION:
                                                                              NAME/KEY: misc_feature
LOCATION: (1)..(24)
                                                                                                                  FEATURE
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OTHER INFORMATION: Viral
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Ryan, Patricia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hallenbeck, Paul
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Forry-Schaudies,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Clarke, Lori
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Jakubczak, John
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                                                               Fig.
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100.0%; Pred. No.
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                                                               SV40 early Poly(A)
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APPLICANT: NICOlaides, Nicholas
APPLICANT: Sass, Philip
APPLICANT: Grasso, Luigi
APPLICANT: Grasso, Luigi
APPLICANT: Kinzler, Kenneth
APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: A METHOD FOR GENERATING HYPERMUTABLE
TITLE OF INVENTION: ORGANISMS
FILE REFERENCE: 01107.00138
CURRENT APPLICATION NUMBER: US/09/853,646
CURRENT FILING DATE: 2000-05-17
PRIOR APPLICATION NUMBER: 60/204,769
PRIOR APPLICATION NUMBER: 60/204,769
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US-09-853-646-4
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                                                                                  ; OTHER INFORMATION: Recombinant US-09-853-646-4
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4
LENGTH: 25
           Matches
                            Query Match
Best Local (
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Best Local Similarity 100.
Than 19; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 4, Application US/09853646
Patent No. US20020055106A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn version 3.1
SEQ ID NO 1
LENGTH: 24
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Publication No. US20030190633A1
GENERAL INFORMATION:
APPLICANT: TANGA, Michifumi
APPLICANT: OKAMURA, Hiroshi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local
Matches :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: OKAMURA, HITCHELD
APPLICANT: TAKAGI, Kenichi
TITLE OF INVENTION: SUPPORT FOR FIXING NUCLEOTIDE AND PROCESS FOR PRODUCING THE SAME
FILE REFERENCE: TANGGA
CURRENT FILING NOBER: US/10/182,434
CURRENT FILING DATE: 2002-07-29
FRIOR APPLICATION NUMBER: UF 2000-019301
PRIOR FILING DATE: 2000-01-27
NUMBER OF SEQ ID NOS: 2
                                                                                                                       ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                           TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
         Local Similarity nes 19; Conserv
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19; Conserv
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         Conservative
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                        100.0%;
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100.0%; Pred. No. 1.7
tive 0; Mismatches
0.7%; Score 19; DB 9; L
100.0%; Pred. No. 1.7e+02;
ive 0; Mismatches 0;
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                                        Length 25;
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     Indels
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                                                                                          Matches
                                                                                                                          Query Match
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Best Local Similarity
Matches 19; Conserv
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SEQ ID NO 52
LENGTH: 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Risinger, Carl APPLICANT: Andersson, Mar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Risinger, Carl
APPLICANT: Andersson, Ma
APPLICANT: Lewander, Tom
                                                                                                                                                                                                 TYPE: DNA
ORGANISM: artificial sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA ORGANISM: artificial sequence FEATURE:
                                                                                                         Local
                                          2661 CAAAAAAAAAAAAAAAA 2679
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                                                                                                         Similarity
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Lewander, Tommy
Olaisson, Erik
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                                                                                  0.7%; Scilarity 100.0%; F
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APPLICANT: Olaisson, Erik
I TITLE OF INVENTION: Detection of CYP2D6 Polymorphisms
FILE REFERENCE: GG119.1US
CURRENT APPLICATION NUMBER: US/09/942,310
CURRENT FILING DATE: 2001-08-29
PRIOR APPLICATION NUMBER: GB 0021286.0
PRIOR FILING DATE: 2000-08-30
I PRIOR FILING DATE: 2000-08-30
I PRIOR FILING DATE: 7000-08-30
I NUMBER OF SEQ ID NOS: 77
SOFTWARE: Patentin version 3.1
SEQ ID NO 59
LENGTH: 25
TYPE: NNA
                                                                                           ; OTHER INFORMATION: synthetic oligonucleotide US-09-942-310-59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Andersson, Maria K.
APPLICANT: Lewander, Tommy
APPLICANT: Olaisson, Erik
TITLE OF INVENTION: Detection of CYP2D6 Polymorphisms
FILE REFERENCE: GG119.1US
CURRENT APPLICATION NUMBER: US/09/942,310
CURRENT FILING DATE: 2001-08-29
PRIOR APPLICATION NUMBER: GB 0021286.0
PRIOR FILING DATE: 2000-08-30
NUMBER OF SEQ ID NOS: 77
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100.0%; Pred. No.
                         Score 19; DB 10;
; Pred. No. 1.7e+02;
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    Mismatches
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                                           Length 25;
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RESULT 17
US-09-099-823-14/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 16
US-10-002-536A-5/c
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Best Local S
Matches 19
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APPLICANT: Nagel, Aaron C.
APPLICANT: Nagel, Aaron C.
APPLICANT: Dombkowski, Alan A.
TITLE OF INVENTION: COMPOSITIONS AND SYSTEMS FOR IDENTIFYING AND COMPARING EXPRESSED TITLE OF INVENTION: (MRNAS) IN EUKARYOTIC ORGANISMS
FILE REFERENCE: 65446-87
CURRENT APPLICATION NUMBER: US/10/002,536A
CURRENT FILING DATE: 2003-02-11
NUMBER OF SEQ ID NOS: 5
SOFTWARE: Patentin version 3.1
SEQ ID NO 5
LENGTH: 25
LENGTH: 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: BILLIN
                                                                                                                                                                                                                                                                                                                            APPLICANT: SCHEFFEH, C. APPLICANT: STROUPE, STEPHEN D.
APPLICANT: YU, HONG
APPLICANT: YU, HONG
TITLE OF INVENTION: REAGENTS AND METHODS USEFUL
TITLE OF INVENTION: FOR DETECTING DISEASES OF THE BREAST
TITLE OF GROUENCES: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
APPLICATION NUMBER: US/09/099,823
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/879,354
FILING DATE: 20-JUN-1997
                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Abbott Laboratories
STREET: 100 Abbott Park Road
                                                                                                              OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
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nes 19; Conserv
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STATE: IL
                                                                                                                                                                                                                                 ZIP: 60064-3500
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COLPITTS, TRACEY L.
FRIEDMAN, PAULA N.
GORDON, JULIAN
                                                                                                                                                                                                                                                    USA
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HODGES, STEVEN C.
KLASS, MICHAEL R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BILLING-MEDEL, PATRICIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                              RUSSELL, JOHN C.
SCHEFFEL, CHRISTI
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Pred. No.
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1.7e+02;
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; Sequence 2, Application US/10182434
; Publication No. US20030190633A1
; GENERAL INFORMATION:
; APPLICANT: TANGA, Michifumi
; APPLICANT: TAKAGI, Kenichi
; APPLICANT: TAKAGI, Kenichi
; APPLICANT: TAKAGI, Kojiro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Grasso, Luigi
APPLICANT: Kinzler, Kenneth
APPLICANT: Kinzler, Kenneth
APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: A METHOD FOR GENERATING HYPERMUTABLE
TITLE OF INVENTION: ORGANISMS
FILE REFERENCE: 01107.00138
CURRENT APPLICATION NUMBER: US/09/853,646
CURRENT APPLICATION NUMBER: 60/204,769
PRIOR FILING DATE: 2001-05-14
PRIOR APPLICATION NUMBER: 60/204,769
PRIOR FILING DATE: 2000-05-17
NUMBER OF SEQ ID NOS: 4
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US-10-182-434-2
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Best Local Similarity
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Matches :
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APPLICANT: OKAMURA, Hiroshi
APPLICANT: TAKAGI, Kenichi
APPLICANT: TAKAHASHI, Kojiro
TITLE OF INVENTION: SUPPORT FOR FIXING NUCLEOTIDE AND PROCESS FOR PRODUCING THE SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Nicolaides, Nicholas
APPLICANT: Sass, Philip
APPLICANT: Grasso, Luigi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Recombinant DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Becker, Cheryl REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 847/938-2623
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                                                                                                                                                                                                                                                                                 2661 САЛАЛАЛАЛАЛАЛАЛА 2679
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100.0%; Pred. No.
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100.0%; Pred. No.
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No.
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CURRENT APPLICATION NUMBER: US/10/182,434

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APPLICANT: Weiner, George
APPLICANT: Hartmann, Gunther
TITLE OF INVENTION: Methods for Enhancing Antibody-Induced
TITLE OF INVENTION: Cell Lysis and Treating Cancer
FILE REFERENCE: C1039/7052 (AWS)
CURRENT APPLICATION NUMBER: US/09/888,326
CURRENT FILING DATE: 2001-06-22
PRIOR APPLICATION NUMBER: US 60/213,346
PRIOR FILING DATE: 2000-06-22
NUMBER OF SEQ ID NOS: 848
SOFTWARE: FASTSEQ for Windows Version 3.0
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US-09-888-326-837/c
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US-09-809-545A-84/c
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                                                           SEQ ID NO 837
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Best Local Similarity 100.0%;
Matches 19; Conservative (
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LENGTH: 27
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TITLE OF INVENTION: SECRETED FACTORS
FILE REFERENCE: SCIOS.017A
CURRENT APPLICATION NUMBER: US/09/809,545A
CURRENT FILING DATE: 2001-03-14
NUMBER OF SEQ ID NOS: 84
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PRIOR APPLICATION NUMBER: JP 2000-019301
PRIOR FILING DATE: 2000-01-27
NUMBER OF SEQ ID NOS: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Stanton, Lawrence W. APPLICANT: White, R. Tyler
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Oligos corresponding to polylinker sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn version 3.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
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Pred. No.
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. 1.7e+02;
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APPLICANT: Petersen, Deanna M.

APPLICANT: Petersen, Deanna M.

APPLICANT: Pouron, Yves

ITITLE OF INVENTION: Immunostimulatory Nucleic Acids for the

ITITLE OF INVENTION: Treatment of Asthma and Allergy

FILE REFERENCE: C1037/7013 (HCL/MAT)

CURRENT APPLICATION NUMBER: US/09/776,479

CURRENT FILING DATE: 2001-02-02

PRIOR APPLICATION NUMBER: US 60/179,991

PRIOR APPLICATION NUMBER: US 60/179,991

PRIOR APPLICATION NUMBER: US 60/179,991

PRIOR FILING DATE: 2000-02-03

NUMBER OF SEQ ID NOS: 1093

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 913

LENGTH: 18

TYPE: DNA

TYPE: DNA

TYPE: DNA
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US-09-776-479-913/c
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                     Sequence 913, Application US/09776479 Publication No. US20030087848A1
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SEQ ID NO 6
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APPLICANT: Roth, Ma
APPLICANT: Lizardi,
APPLICANT: Feng, L
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Publication No. US20030082556A1
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PRIOR FILING DATE: 2000-08-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/994,311
CURRENT FILING DATE: 2001-11-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Binary Encoded Sequence Tags FILE REFERENCE: AGL 100
                                     ORGANISM: Artificial Sequence
OTHER INFORMATION: Synthetic Sequence
                             FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
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LOCATION: (0)...(0)
OTHER INFORMATION: phosphorothioate backbone
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18; Conservative 0;
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Latimer, Darin R.
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Lizardi, Paul M.
Fenz
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100.0%; Pred. No.
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Pred. No.
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US-09-776-479-913

Query Match

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DB 10; 1

Length 18;

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US-09-370-541-14/c
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LENGTH: 18
TYPE: DNA
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APPLICANT: Manchazan, Muthiah
APPLICANT: Cook, Phillip Dan
APPLICANT: Prakash, Thazha P
APPLICANT: Kawasaki, Andrew M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 939, Application US/09776479 Publication No. US20030087848A1 GENERAL INFORMATION:
                                                                SOFTWARE: PatentIn Ver. SEQ ID NO 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 14, Application US/09370541 Publication No. US20030088079A1
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APPLICANT: Fouron, Yves
TITLE OF INVENTION: Immunostimulatory Nucleic Acids for the
TITLE OF INVENTION: Treatment of Asthma and Allergy
FILE REFERENCE: C1037/7013 (HCL/MAT)
CURRENT APPLICATION NUMBER: US/09/776,479
CURRENT FILING DATE: 2001-02-02
PRIOR APPLICATION NUMBER: US 60/179,991
PRIOR FILING DATE: 2000-02-03
NUMBER OF SEQ ID NOS: 1093
SOFTWARE: FASTSEQ for Windows Version 3.0
                                                                                                                                              EARLIER APPLICATION NUMBER: 09/130,973
EARLIER FILING DATE: 1998-08-07
EARLIER APPLICATION NUMBER: 09/016,520
EARLIER FILING DATE: 1998-01-30
EARLIER APPLICATION NUMBER: 60/037,143
EARLIER FILING DATE: 1997-02-14
EARLIER APPLICATION NUMBER: 09/344,260
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APPLICANT: Petersen, Deanna M
APPLICANT: Fouron, Yves
                                                                                                        EARLIER FILING DATE: 1999-06-25
NUMBER OF SEQ ID NOS: 21
                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/370,541
CURRENT FILING DATE: 1999-08-09
                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Aminoxxy-Modified Nucleosidic Compounds And Oligomeric TITLE OF INVENTION: Compounds Prepared Therefrom
                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: ISIS399:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Artificial Sequence FEATURE:
OTHER INFORMATION: Synthetic Sequence
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity 100.
les 18; Conservative
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18; Conserv
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100.0%; Pr
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100.0%; Pred. No.
tive 0; Mismatcl
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; OTHER INFORMATION: Description of Artificial Sequence: antisense; OTHER INFORMATION: sequence
US-09-370*541-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Bratzler, Robert L.
APPLICANT: Petersen, Deanna M.
APPLICANT: Petersen, Deanna M.
APPLICANT: Pouron, Yves
TITLE OF INVENTION: Immunostimulatory Nucleic Acids for
TITLE OF INVENTION: Treatment of Asthma and Allergy
FILE REFERENCE: C1037/7013 (HCL/MAT)
CURRENT APPLICATION NUMBER: US/09/776,479
CURRENT FILING DATE: 2001-02-02
PRIOR APPLICATION NUMBER: US 60/179,991
PRIOR FILING DATE: 2000-02-03
NUMBER OF SEQ ID NOS: 1093
SOFTWARE: FastSEQ for Windows Version 3.0
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                                                              SEQ ID NO 939
LENGTH: 18
TYPE: DNA
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Matches
                                                                                                                                                                                                                                                                                                                                                                                     Publication No. US20040067902A9 GENERAL INFORMATION:
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TITLE OF INVENTION: Immunostimulatory Nucleic Acids for the
TITLE OF INVENTION: Treatment of Asthma and Allergy
FILE REFERENCE: C1037/7013 (HCL/MAT)
CURRENT APPLICATION NUMBER: US/09/776,479
CURRENT FILING DATE: 2001-02-02
                                                                                                                          PRIOR APPLICATION NUMBER: US 60/179,991
PRIOR FILING DATE: 2000-02-03
NUMBER OF SEQ ID NOS: 1093
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                       APPLICANT: Bratzler, Robert L.
APPLICANT: Petersen, Deanna M.
APPLICANT: Fouron, Yves
OTHER INFORMATION: Synthetic Sequence
                    ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 18
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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Local Similarity 100.0%;
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Pred. No.
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hes 0;
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RESULT 29
US-10-125-295-9/c
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US-09-979-275A-7/c
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Matches 18; Conservative (
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SEQ ID NO 7
LENGTH: 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: NACAL, HIROSHI
APPLICANT: NACAL, KYOKO
APPLICANT: NAKAJIMA, TERUMI
TITLE OF INVENTION: NOVEL PROTEINS HAVING HEMOLYTIC ACTIVITY AND GENES
FILE OF INVENTION: ENCODING THE PROTEIN
FILE REFERENCE: 037181.50611US
CURRENT APPLICATION NUMBER: US/09/979,275A
CURRENT FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: PCT/JP01/02209
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: JP 2000-78967
RIOR FILING DATE: 2000-03-21
PRIOR FILING DATE: 2000-03-21
NUMBER OF SEQ ID NOS: 12
COUNTY NEWS APPLICATION NUMBER: JP 2000-78967
NUMBER OF SEQ ID NOS: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: this sequence may encompass 12-18 nucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                     Wallace, Robert Bruce
Cossman, Jeffrey
French, Cynthia
TITLE OF INVENTION: Lyophilization of Cultured Human Cells
to Preserve RNA and DNA
                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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                                                                                                                      COUNTRY: USA
ZIP: 94111-3834
                                                                                                                                                                  STATE: California
                                                                                                                                                                                       CITY: San Francisco
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APPLICATION DATA:
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100.0%; Pred. No. 5.4
ive 0; Mismatches
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100.0%; Pred. No. 5.4
ive 0; Mismatches
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hes 0;
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APPLICANT: Kohse, Peter
APPLICANT: Hohse, Peter
APPLICANT: Magner, Richard
TITLE OF INVENTION: Peptide Acceptor Ligation Methods
FILE REFERENCE: 50036/031002
CURRENT APPLICATION NUMBER: US/10/208,357
CURRENT FILING DATE: 2002-07-30
PRIOR APPLICATION NUMBER: US/09/619,103
PRIOR APPLICATION NUMBER: US/09/619,103
PRIOR FILING DATE: 1099-07-19
PRIOR FILING DATE: 1099-07-27
NUMBER OF SEO ID NOS: 26
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US-10-208-357-24
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ENGTH: 18
                                                                                  Matches
                                                                                                                    Query Match
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Best Local Similarity
Matches 18; Conserv
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Publication No. US20020182687A1
                                                                                                                                                           FEATURE: OTHER INFORMATION: designed sequence for nucleic acid purification
                                                                                                                                                                                                           ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                            TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 9: SEQUENCE CHARACTERISTICS:
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FILING DATE: 17-Apr-2002
CLASSIFICATION: cUtchown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/545,225
FILING DATE: 07-Apr-2000
APPLICATION NUMBER: US 08/884,029
FILING DATE: 27-JUN-1997
ATTORNEY/AGENT INFORMATION:
                                                                             Local Similarity es 18; Conserv
                        2662 AAAAAAAAAAAAAAAAAA 2679
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LOCATION: 13..18
OTHER INFORMATION: /mod base= OTHER
/note= "t at positions 13-18 may be
present or absent"
SEQUENCE DESCRIPTION: SEQ ID NO: 9:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Parent, Annette S.
REGISTRATION NUMBER: 42,058
REFERENCE/DOCKET NUMBER: 02558B-059100US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2662 АЛАЛАЛАЛАЛАЛАЛ 2679
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AAAAAAAAAAAAAAA 18
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                                                                         0.7%; So illarity 100.0%; F Conservative 0;
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100.0%; Pred. No. 5.4e+02;
tive 0; Mismatches 0;
                                                                       Score 18; DB 13;
; Pred. No. 5.4e+02;
0; Mismatches 0;
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APPLICANT: Bratzler, Robert L.
TITLE OF INVENTION: Inhibition of Angiogenesis by
FILE REFERENCE: C1037/7025 (HCL/MAT)
CURRENT APPLICATION NUMBER: US/10/017,995
CURRENT FILING DATE: 2001-12-18
PRIOR APPLICATION NUMBER: US 60/255,534
PRIOR FILING DATE: 2000-12-14
NUMBER OF SEQ ID NOS: 1093
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 913
LENGTH: 18
TYPE: DNA
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US-10-017-995-939/c
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US-10-017-995-913/c
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US-10-112-653-882/c
                                                                                                                                                                                                                                                                                                                                                                                                                      ; OTHER INFORMATION: Synthetic Sequence US-10-017-995-913
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Sequence 939, Application US/10017995
Publication No. US20030055014A1
GENERAL INFORMATION:
APPLICANT: Bratzler, Robert L.
TITLE OF INVENTION: Inhibition of Angiogenesis by Nucleic Acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 913, Application US/10017995 Publication No. US20030055014A1 GENERAL INFORMATION:
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SEQ ID NO 882
LENGTH: 18
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Publication No. US20030050268A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.7%; Score 18; DB Best Local Similarity 100.0%; Pred. No. 5.4 Matches 18; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                   Query Match 0.7%; Score 18; DB 14; Length 18; Best Local Similarity 100.0%; Pred. No. 5.4e+02; Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Krieg, Arthur M.
APPLICANT: Berg, Daniel J.
APPLICANT: Berg, Daniel J.
TITLE OF INVENTION: IMMUNOSTIMULATORY NUCLEIC ACID FOR
TITLE OF INVENTION: TREATMENT OF NON-ALLERGIC INFLAMMATORY DISEASES
FILE REFERENCE: C01039/70060(AWS)
CURRENT APPLICATION NUMBER: US/10/112,653
CURRENT FILING DATE: 2002-03-29
PRIOR APPLICATION NUMBER: US 60/279,642
PRIOR FILING DATE: 2001-03-29
NUMBER OF SEQ ID NOS: 1040
NUMBER OF SEQ ID NOS: 1040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Synthetic Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
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                                                                                                                                                                                                                                                                 2662 AAAAAAAAAAAAAAAAA 2679
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           by Nucleic Acids
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US-10-056-479A-15/c

Sequence 15, Application US/10056479A Publication No. US20030175678A1 GENERAL INFORMATION:

APPLICANT: Bowen, Banjamin A.
APPLICANT: Deakin, Edward
APPLICANT: Goldsmith, Neil
APPLICANT: Haudenschild, Christian
APPLICANT: Houck, David

APPLICANT:

McAlpine, James B. Neilsen, Soren Pazoles, Christopher RESULT 35

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PRIOR FILING DATE: 2000-12-14
NUMBER OF SEQ ID NOS: 1093
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 939
LENGTH: 18
TYPE: DNA
                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Xu, APPLICANT: Jab. APPLICANT: You APPLICANT: He, APPLICANT: The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 4, Application US/10206613
Publication No. US20030104432A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity 100.0%; Matches 18; Conservative (
                                                                      Matches
                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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APPLICANT: He, Biao
APPLICANT: The Regents of the University of California
APPLICANT: The Regents of the University of California
TITLE OF INVENTION: Methods of Amplifying Long Sense Sti
FILE REFERENCE: 023070-119510US
CURRENT APPLICATION NUMBER: US/10/206,613
CURRENT FILING DATE: 2002-11-04
CURRENT FILING DATE: 2002-11-04
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CURRENT APPLICATION NUMBER: US/10/017,995
CURRENT FILING DATE: 2001-12-18
                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 60/308,190 PRIOR FILING DATE: 2001-07-27 NUMBER OF SEQ ID NOS: 8
                                                                                                                                                              OTHER INFORMATION:
                                                                                                                                                                                                LENGTH: 18
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: Synthetic Sequence
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                                  2662 AAAAAAAAAAAAAAAAA 2679
18 AAAAAAAAAAAAAAAA 1
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Jablons, David
                                                                      Conservative
                                                                                                                                                            Description of Artificial Sequence:oligo dT-18 linker primer
                                                              100.0%; Pi
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                                                                                     0.7%; Score 18;
100.0%; Pred. No.
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100.0%; Pred. No.
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                                                                        Mismatches
                                                                                       DB 15; 1
5.4e+02;
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CURRENT APPLICATION NUMBER: US/10/056,479A
CURRENT FILING DATE: 2003-02-07
PRIOR APPLICATION NUMBER: US 60/263,807
PRIOR FILING DATE: 2001-01-24
NUMBER OF SEO ID NOS: 15
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 15
LENGTH: 18
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE: TIPE: DNA
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US-10-352-704-12/c
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Best Local Similarity
Matches 18; Conserv
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                                                           TELEPHONE: (202)638-6666
TELEPAX: (202) 393-5350
TELEX: RCA 248593 IDEA UR
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Spencer, Marget E.
APPLICANT: Stafford, Angela
TITLE OF INVENTION: Methods for Identifying Genes Regulating
TITLE OF INVENTION: Desired Cell Phenotypes
FILE REFERENCE: 50273/005002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                   APPLICATION NUMBER: US/08/358,556A
FILING DATE: 14-DEC-1994
APPLICATION NUMBER: FR 9315164
FILING DATE: 16-DEC-1993
ATTORNEY/AGENT INFORMATION:
NAME: Player, William E.
REGISTRATION NUMBER: 31,409
REFERENCE/DOCKET NUMBER: 10577/P58418
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/10/352,704
FILING DATE: 28-Jan-2003
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Process for Preparing Polynucleotides on a Solid Support and Apparatus Permitting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2662 AAAAAAAAAAAAAAAAAA 2679
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Chatelain, Francois
Kumarev, Viktor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18 АААЛААААААААААА 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Washington D.C STATE: D.C
STRANDEDNESS: single
                  LENGTH: 18 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Jacobson, Price, Holman & Stern STREET: 400 Seventh St. N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.7%; Score 18; DB 15; ilarity 100.0%; Pred. No. 5.4e+02; Conservative 0; Mismatches 0;
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                                                                                                                                                                                                    TELEX: RCA 248593 IDEA UR INFORMATION FOR SEQ ID NO: 18:
                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/358,556A
FILING DATE: 14-DEC-1994
APPLICATION NUMBER: FR 9315164
FILING DATE: 16-DEC-1993
ATTORNEY/AGENT INFORMATION:
NAME: Player, William E.
REGISTRATION NUMBER: 31,409
REGISTRATION NUMBER: 10577/P58418
TELEPHONE: (202) 393-5350
TELEPHONE: (202) 393-5350
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/352,704
FILING DATE: 28-Jan-2003
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                    FEATURE:
                                  ANTI-SENSE: NO FRAGMENT TYPE: N-terminal
                                                                      STRANDEDNESS: single TOPOLOGY: linear MOLECULE TYPE: DNA (genomi HYPOTHETICAL: NO
                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 31
CORRESPONDENCE ADDRESS:
ADDRESSE: Jacobson, Price, Holman & Stern
STREET: 400 Seventh St. N.W.
CITY: Washington D.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kumarev, Viktor
TITLE OF INVENTION: Process for Preparing Polynucleotides on a Solid Support and Apparatus Permitting Implementation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: CDS
LOCATION: 1..18
SEQUENCE DESCRIPTION: SEQ ID NO: 12:
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MOLECULE TYPE: DNA (genomic)
HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: N-terminal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Chatelain, Francois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2662 АЛАЛАЛАЛАЛАЛАЛА 2679
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                                                                                                                                         TYPE: nucleic acid
 NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 20004
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                                                                                         DNA (genomic)
CDS
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; Pred. No. 5.4e+02;
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Gaps

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; SEQUENCE DESCRIPTION: SEQ ID NO: 18: US-10-352-704-18
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US-10-075-335-9/c
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                                                                                                                                                                             APPLICANT: BEDIAN, VAHE
APPLICANT: GLADUE, RONALD F
APPLICANT: CORVALAN, JOSE
APPLICANT: JIA, XIAO-CHI
APPLICANT: FENG, XIAO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Loc
Matches
                                                                                                                                                                                                                                                                                       Sequence 144, Application US/10 Publication No. US20030211100A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 10
SOFTWARE: PatentIn version 3.1
SEQ ID NO 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 9, Application US/10075335 Publication No. US20030186237A1 GENERAL INFORMATION:
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SOFTWARE: PatentIn Ver. SEQ ID NO 144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/10/075,335
CURRENT FILING DATE: 2003-01-08
PRIOR APPLICATION NUMBER: 60/268,664
PRIOR FILING DATE: 2001-02-14
PRIOR APPLICATION NUMBER: 60/348,242
PRIOR FILING DATE: 2001-11-07
PRIOR FILING DATE: 2001-11-07
                                  TITLE OF INVENTION: ANTIBODIES TO CD40
FILE REFERENCE: ABX-PF/3 US
CURRENT APPLICATION NUMBER: US/10/292,088
CURRENT FILING DATE: 2003-03-14
PRIOR APPLICATION NUMBER: 60/348,980
PRIOR FILING DATE: 2001-11-09
RUMBER OF SEQ ID NOS: 147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/268,645
PRIOR FILING DATE: 2001-02-14
PRIOR PPLICATION NUMBER: 60/344,557
PRIOR FILING DATE: 2001-11-07
PRIOR APPLICATION NUMBER: 60/306,216
PRIOR FILING DATE: 2001-07-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Che, Shaoli
TITLE OF INVENTION: Methods and Compositions of Amplifying RNA
FILE REFERENCE: HO-P02202US2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/350,176
PRIOR FILING DATE: 2001-11-09
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ORGANISM: Artificial Sequence
FEATURE:
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nes 18; Conservative
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nes 18; Conserv
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100.0%; Pred. No.
tive 0; Mismatch
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. 5.4e+02;
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5.4e+02;
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PRIOR FILING DATE: 1999-09-25
PRIOR APPLICATION NUMBER: US 60/156,135
PRIOR FILING DATE: 1999-09-27
PRIOR APPLICATION NUMBER: US 60/227,436
PRIOR FILING DATE: 2000-08-23
NUMBER OF SEQ ID NOS: 1145
SOPTWARE: FASISEQ for Windows Version 3.0
SEQ ID NO 913
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US-10-314-578-939/c
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US-10-314-578-913/c
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Best Local Similarity 100.
Matches 18; Conservative
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                                                                                                                                                                                                                                                                                                        Sequence 939, Application US/10314578 Publication No. US20030212026A1
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                                                                                          APPLICANT: Schetter, Christian
APPLICANT: Vollner, Jorg
TITLE OF INVENTION: Immunostimulatory Nucleic Acids
FILE REFERENCE: C1039/7035 (HCL/MAT)
CURRENT APPLICATION NUMBER: US/10/314,578
CURRENT FILING DATE: 2002-12-09
PRIOR APPLICATION NUMBER: US 60/156,113
PRIOR PILING DATE: 1999-09-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Krieg, Arthur M.
APPLICANT: Schetter, Christian
APPLICANT: Vollmer, Jorg
TITLE OF INVENTION: Immunostimulatory Nucleic Acids
FILE REFERENCE: C1039/7035 (HCL/MAT)
CURRENT APPLICATION NUMBER: US/10/314,578
CURRENT FILING DATE: 2002-12-09
          PRIOR APPLICATION NUMBER: US 60/156,135
PRIOR FILING DATE: 1999-09-27
PRIOR APPLICATION NUMBER: US 60/227,436
PRIOR FILING DATE: 2000-08-23
                                                                                                                                                                                                                                           APPLICANT: Krieg, Arthur M. APPLICANT: Schetter, Chris
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TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Synthetic Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.7%; Score 18; DB 15; 100.0%; Pred. No. 5.4e+02;
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100.0%; Pred. No.
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RESULT 42
US-10-389-155-97/c
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                                                                                             TELEPAX: (415) 576-0
INFORMATION FOR SEQ ID NO: 97:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: FastSEQ for Windows Version 3.0 SQ ID NO 939
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 100.0%; 105 18; Conservative 0
                                                                                                                                                                                                                                   APPLICATION NUMBER: US 08/484,537 FILING DATE: 07-JUN-1995 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/389,155
              MOLECULE
                                                                                                                                                             REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR
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R APPLICATION DATA:
APPLICATION NUMBER: US/09/325,000

FILING DATE: 01-UN-1999

APPLICATION NUMBER: US 07/290,975

FILING DATE: 28-DEC-1988
                                         TYPE: nucleic acid
STRANDEDNESS: single
                                                                            LENGTH: 18 base pairs
                                                                                                                                                                                             REGISTRATION NUMBER: 30,223
                                                                                                                                               TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 07/310,252
FILING DATE: 13-FEB-1989
APPLICATION NUMBER: US 07/590,274
FILING DATE: 28-SEP-1990
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                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US 07/634,278 FILING DATE: 19-DEC-1990
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                              ÖGY:
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Schneider, William P.
Landolfi, Nicholas F.
Coelingh, Kathleen L.
Selick, Harold E.
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            DNA
                            linear
                                                                                                                                                                                                                 William M
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Pred. No.
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5.4e+02;
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LENGTH: 18
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PRIOR APPLICATION NUMBER: 60/343,169
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APPLICANT: Ghazala Hashm
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                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
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; OTHER INFORMATION: Probe sequence derived from human genomic US-10-271-602B-84
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PRIOR FILING DATE: 2001-10-14
PRIOR PELICATION NUMBER: 60/329,620
PRIOR FILING DATE: 2001-10-15
PRIOR APPLICATION NUMBER: 60/329,428
PRIOR APPLICATION NUMBER: 60/329,428
PRIOR FILING DATE: 2001-10-14
PRIOR APPLICATION NUMBER: 60/329,619
PRIOR FILING DATE: 2001-10-15
PRIOR PILING DATE: 2001-20-14
PRIOR PILING DATE: 2002-03-14
NUMBER OF SEQ ID NOS: 212
                                                                                                                                                                        Sequence 204, Application US/10334143 Publication No. US20040009549A1 GENERAL INFORMATION:
APPLICANT: SUDARSANAM, SUCHA
TITLE OF INVENTION: METHOD FOR DETECTING REMOTE HOMOLOGUES AND NOVEL
TITLE OF INVENTION: KINASES IDENTIFIED WITH THE METHOD
FILE REFERENCE: 038602/1543
CURRENT APPLICATION NUMBER: US/10/334,143
CURRENT FILING DATE: 2002-12-31
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                                                                                                                          APPLICANT: GRIGORIEV, IGOR VYACHESLAVOVICH APPLICANT: SUDARSANAM, SUCHA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Michael Seul
TITLE OF INVENTION: MULTIPLEXED ANALYSIS OF POLYMORPHIC LOCI
TITLE OF INVENTION: BY CONCURRENT INTERROGATION AND ENZYME-MEDIATED DETECTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: eMAP-US
CURRENT APPLICATION NUMBER: US/10/271,602B
CURRENT FILING DATE: 2002-10-15
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                                                                                                                                                                                                                                                                                                                                             AAAAAAAAAAAAAAAA 18
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OTHER INFORMATION: /mod
                                                                                                                                                                                                                                                                                                                                                                                                                                         0.7%; Score 18; DB 16;
llarity 100.0%; Pred. No. 5.4e+02;
Conservative 0; Mismatches 0;
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100.0%; Pr
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5.4e+02;
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NUMBER OF SEQ ID NOS: 207
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 204
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INFORMATION
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                                               APPLICATION NUMBER: US 07/290,975
PILING DATE: 28-DEC-1988
APPLICATION NUMBER: US 07/310,252
PILING DATE: 13-FEB-1989
APPLICATION NUMBER: US 07/590,274
FILING DATE: 28-SEP-1990
APPLICATION NUMBER: US 07/634,278
FILING DATE: 19-DEC-1990
APPLICATION NUMBER: US 08/484,537
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 011823-0026
TELECOMMUNICATION INFORMATION:
FELEPHONE: (415) 736-030
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER RADABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC Compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Improved Humanized Immunoglobulins NUMBER OF SEQUENCES: 100
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                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/10/389,417 FILING DATE: 13-Mar-2003 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/09/325,000 FILING DATE: 01-JUN-1999
                 TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Application US/10389417
No. US20040049014A1
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Schneider, William P.
Landolfi, Nicholas F.
Coelingh, Kathleen L.
Selick, Harold E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Queen, Cary L.
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                                                                              011823-002650US
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GENERAL

INFORMATION:

APPLICANT: Bowen, Banjamin A. APPLICANT: Deakin, Edward APPLICANT: Goldsmith, Neil APPLICANT: Haudenschild, Chr

Goldsmith, Neil Haudenschild, Christian Houck, David

Pazoles, Christopher McAlpine, James B.

Soren

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                                                      RESULT 47
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                                 US-10-785-744-15/c
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Best Local Sim:
Matches 18;
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     Sequence 15, Ap
Publication No.
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                                                                                                                                                                          Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 26, Ap
Publication No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: COMPOSITIONS AND METHODS FILE REFERENCE: 3826-0011 CURRENT APPLICATION NUMBER: US/10/653,416 CURRENT FILING DATE: 2003-09-03 PRIOR APPLICATION NUMBER: 60/407,248 PRIOR FILING DATE: 2002-09-03 NUMBER OF SEQ ID NOS: 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: RASHTCHIAN, AYOUB APPLICANT: SCHUSTER, DAVID M.
                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Ver. 3.2
                                                                                                                                                                                                                                                           OTHER INFORMATION: OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: oligonucleotide
                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 18
TYPE: DNA
                                                                                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                                                              FEATURE:
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                                                                                                                                      2662 АЛЛАЛАЛАЛАЛАЛАЛА 2679
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TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                         Similarity
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LOCATION: 13..18
OTHER INFORMATION: /mod_base= OTHER
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LOCATION:
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Application US/10785744
No. US20040133941A1
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No. US20040110201A1
                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                                           this sequence may encompass 12-18 nucleotides according to the specification as filed
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100.0%; Pred. No.
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100.0%; Pred. No. 5.4
ive 0; Mismatches
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                                                                                                                                                                      DB 17; I
5.4e+02;
hes 0;
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US-10-628-525-30
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US-10-735-592-1/c
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                                                                                                                                                         Sequence 30, Application US/10628525
Publication No. US20040185114A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Art, Krieg
APPLICANT: Joerg, Vollmer
TITLE OF INVENTION: 5' CPG Nucleic Acids and Methods of Use
FILE REFERENCE: C1037.70038US01
CURRENT APPLICATION NUMBER: US/10/735,592
CURRENT FILING DATE: 2003-12-11
NUMBER OF SEQ ID NOS: 69
NUMBER OF SEQ ID NOS: 69
SOFTWARE: PatentIn version 3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Methods for Identifying Genes Regulating TITLE OF INVENTION: Desired Cell Phenotypes FILE REFERENCE: 50273/005002 CURRENT EPPLICATION NUMBER: US/10/785,744 CURRENT FILING DATE: 2004-02-23 FRIOR APPLICATION NUMBER: US/10/056,479 PRIOR PILING DATE: 2003-02-07 PRIOR FILING DATE: 2003-02-07 PRIOR PILING DATE: 2003-02-07 PRIOR APPLICATION NUMBER: US 60/263,807 PRIOR FILING DATE: 2001-01-24 NUMBER OF SEQ ID NOS: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ENGTH: 18
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Guan, Hanping
TITLE OF INVENTION: Starch Encapsulation
NUMBER OF SEQUENCES: 37
CORRESPONDENCE ADDRESS:
ADDRESSEE: Greenlee, Winner and Sullivan, P.C.
STREET: 5370 Manhattan Circle
                                                                                                                                                                                                                                                                                                                                             2662 AAAAAAAAAAAAAAAAA 2679
                                                                                                                                  APPLICANT: Keeling, Peter
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Stafford, Angela
                                                                                                                                                                                                                                                                                                                                                                                                                        0.7%; Score 18; DB 17; 100.0%; Pred. No. 5.4e+0;
                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                        5.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 18;
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US-10-849-072-21
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; Sequence 21, Application US/10849072
; Publication No. US20040214221A1
; GENERAL INFORMATION:
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                                                                                                                                                                                   APPLICANT: Roche Diagnostics GmbH
TITLE OF INVENTION: High density labeling of DNA with modified or
TITLE OF INVENTION: "Chromophore" carrying nucleotides and DNA polymerases
TITLE OF INVENTION: used
FILE REFERENCE: 4780/00/MO
CURRENT APPLICATION NUMBER: US/10/849,072
CURRENT FILING DATE: 2004-05-19
NUMBER OF SEQ ID NOS: 26
SOFTWARB: Patentin Ver. 2.1
LENGTH: 18
    Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 100
Matches 18; Conservative
                                                                                                OTHER INFORMATION: Description of Artificial Sequence: second OTHER INFORMATION: fragment of SEQ ID NO: 6
                                                                                                                                             ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: 30:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC. Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/628,525
FILING DATE: 28-Jul-2003
CLASSIFICATION BO0
PRIOR APPLICATION NUMBER: US/08/941,445
APPLICATION NUMBER: US/08/941,445
FILING DATE: 30-SEP-1997
APPLICATION NUMBER: US/08/941,445
FILING DATE: 30-SEP-1996
ATTORNEY/AGENT INFORMATION:
NAMME. Witness US/08/941
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2662 AAAAAAAAAAAAAAAA 2679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Winner, Ellen P
REGISTRATION NUMBER: 28,547
REFERENCE/DOCKET NUMBER: 89-
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HYPOTHETICAL: NO SEQUENCE DESCRIPTION: SEQ ID NO: 30:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: Not Relevant MOLECULE TYPE: cDNA to mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
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TELEFAX: (303) 4
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    Conservative
                      100.0%;
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100.0%; Pred. No.
                    0.7%; Score 18; DB 18; 100.0%; Pred. No. 5.4e+0;
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499-8089
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    Mismatches
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                      5.4e+02;
                                     Length 18;
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2662 AAAAAAAAAAAAAAAAA 2679

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AAAAAAAAAAAAAAAA 18

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RESULT 53
US-10-831-778-939/c
US-10-831-778-939/c
; Sequence 939, Application US/10831778
; Publication No. US20040235774A1
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FEATURE:
OTHER INFORMATION: Synthetic Sequence
US-10-831-778-913
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US-10-831-778-913/c
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                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: FOURDI, YVES
TITLE OF INVENTION: Immunostimulatory Nucleic Acids for the
TITLE OF INVENTION: Treatment of Asthma and Allergy
FILE REFERENCE: C1037/7013 (HCL/MAT)
CURRENT APPLICATION NUMBER: US/10/831,778
CURRENT FILING DATE: 2004-04-23
PRIOR APPLICATION NUMBER: US 60/179,991
PRIOR FILING DATE: 2000-02-03
NUMBER OF SEQ ID NOS: 1093
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 913
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
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Best Local (
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LENGTH: 18
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TITLE OF INVENTION: High density labeling of DNA with modified or
TITLE OF INVENTION: "chromophore" carrying nucleotides and DNA po
TITLE OF INVENTION: used
FILE REFERENCE: 4780/00/WO
CURRENT APPLICATION NUMBER: US/10/849,072
CURRENT FILING DATE: 2004-05-19
NUMBER OF SEQ ID NOS: 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Bratzler, Robert L. APPLICANT: Petersen, Deanna MAPPLICANT: Fouron, Yves
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OTHER INFORMATION: Description of Artificial Sequence: second
OTHER INFORMATION: fragment of SEQ ID NO: 6
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                                                                                                                                                                                                 0.7%; Score 18;
100.0%; Pred. No.
tive 0; Mismatcl
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APPLICANT: Bratzler, Robert L.
APPLICANT: Petersen, Deanna M.
APPLICANT: Petersen, Deanna M.
APPLICANT: Potersen, Yves
ITILE OF INVENTION: Immunostimulatory Nucleic Acids for the
TITLE OF INVENTION: Treatment of Asthma and Allergy
FILE REFERENCE: C1037/7013 (HCL/MAT)
CURRENT APPLICATION NUMBER: US/10/831,778
CURRENT FILING DATE: 2004-04-23
PRIOR APPLICATION NUMBER: US 60/179,991
PRIOR APPLICATION STREET CONTROL 10/8
SOFTWARE: FastSEQ for Windows Version 3.0
SOFTWARE: FastSEQ for Windows Version 3.0
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US-10-776-933-150/c
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SEQ ID NO 112
LENGTH: 18
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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                                           APPLICANT: HANSEN, I
APPLICANT: THRUE,
APPLICANT: WESTERGE
APPLICANT: PETERSEE
APPLICANT: WISSENBA
                                                                                                                                             Sequence 150, Applic Publication No. US20 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 112, Application US/10674159A Publication No. US20040242518A1
                                                                                                                                                                                                                                                                                                                                                                                  Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Chen, Jianzhu
APPLICANT: Ge, Qing
APPLICANT: Bisen, Herman
TITLE OF INVENTION: Influenza Therapeutic
FILE REFERENCE: 0492611-0506
CURRENT APPLICATION NUMBER: US/10/674,159A
CURRENT FILING DATE: 2003-09-29
NUMBER OF SEQ ID NOS: 271
APPLICANT: THRUE, CHARLOTTE ALBAEK
APPLICANT: WESSTERGARD, MAJKEN
APPLICANT: PETERSEN, KAMILLE DUMONG
APPLICANT: WISSENBACH, MARGIT
TITLE OF INVENTION: OLIGOMERIC COMPOUNDS FOR THE MODULATION OF THIOREDOXIN
TITLE OF INVENTION: EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 18
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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                                                                                                                                                                     Application US/10776933 o. US20040241717A1
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100.0%; Pro
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100.0%; Pred. No. 5.4e+02;
htive 0; Mismatches 0;
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Pred. No.
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hes 0;
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                                                        US-10-766-096-9/c
                                                                               RESULT 57
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              Sequence 9, Application US/10766096 Publication No. US20040265786A1
                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: DK 2003-01539
PRIOR FILING DATE: 2003-10-20
NUMBER OF SEQ ID NOS: 201
SOFTWARE: Patentin Ver. 3.2
SEQ ID NO 141
LENGTH: 18
TYPE: DNA
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: 58609(71432)
CURRENT APPLICATION NUMBER: US/10/776,917
CURRENT FILING DATE: 2004-02-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: WESTERGAARD, MAJKEN
APPLICANT: PETERSEN, KAMILLE DUMONG
APPLICANT: WISSENBACH, MARGIT
TITLE OF INVENTION: OLIGOMERIC COMPOUNDS FOR THE MODULATION OF RAS EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/446,363 PRIOR FILING DATE: 2003-02-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: HANSEN, BO
APPLICANT: THRUE, CHARLOTTE ALBAEK
APPLICANT: WESTERGAARD, MAJKEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/10/776,933
CURRENT FILING DATE: 2004-02-10
PRIOR APPLICATION NUMBER: 60/446,374
PRIOR FILING DATE: 2003-02-10
NUMBER OF SEQ ID NOS: 150
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
OTHER INFORMATION: poly-T oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial sequence
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ORGANISM: Artificial Sequence
FEATURE:
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les 18; Conserv
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100.0%; Pred. No. 5.4
ive 0; Mismatches
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Pred. No. 5.4e+02;
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ches 0;
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RESULT 58
US-10-872-984-6/c
                                                                                         GENERAL INFORMATION:
APPLICANT: Kaufman,
APPLICANT: Roth, Mai
APPLICANT: Lizardi,
APPLICANT: Feng, Li
                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-10-766-096-9
                                                                                                                                                                                        Sequence 6, Application US/10872984
Publication No. US20040265888A1
                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                   APPLICANT: Feng, Li
APPLICANT: Latimer, Darin R.
TITLE OF INVENTION: Binary Encoded Sequence
FILE REFERENCE: AGL 100
CURRENT APPLICATION NUMBER: US/10/872,984
CURRENT FILING DATE: 2004-06-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: (415) 576-0200
TELEPAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                          2662 АЛАЛАЛАЛАЛАЛАЛАЛА 2679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/884,029
ETILING DATE: 27-UIN-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 02558B-059100US TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/766,096
FILING DATE: 27-Jan-2004
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Lyophilization of Cultured Human Cells to Preserve RNA and DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Lin, Ching-I Patsy
                                                                                                                                                                                                                                                                                                           18 AAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                    18;
                                                                                                                              Roth, Matthew E.
                                                                                                              Roth, Matthew E.
Lizardi, Paul M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: modified_base
LOCATION: 13.18
OTHER INFORMATION: /mod_base= OTHER
/note= "t at positions 13-18 may be
present or absent"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Parent, Annette S. REGISTRATION NUMBER: 42,058
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 94111-3834
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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Cossman, Jeffrey
French, Cynthia
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Pred. No.
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hes 0;
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PRIOR APPLICATION NUMBER: US/09/994,311

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                                                                                                                                                                                                                                                                                               RESULT 60
US-09-917-138-1/c
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US-10-638-141-10/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: Description of Artificial Sequence: Primer US-10-872-984-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 100.0%; F
Marches 18; Conservative 0;
                                                                                                                                                                                            Sequence 1, Application US/09917138
Patent No. US30020031776A1
GENERAL INFORMATION:
APPLICANT: TULLIS, Richard
APPLICANT: STEIFFEL, Jerome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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                                                   TITLE OF INVENTION: ENZYMATIC LABELLING AND DETECTION OF DNA TITLE OF INVENTION: HYBRIDIZATION PROBES FILE REFERENCE: 24730-2207B
CURRENT APPLICATION NUMBER: US/09/917,138
CURRENT FILING DATE: 2001-07-26
PRIOR APPLICATION NUMBER: 09/580,358
PRIOR FILING DATE: 2000-05-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/10/638,141
CURRENT FILING DATE: 2003-08-07
PRIOR APPLICATION NUMBER: US/09/665,728
PRIOR FILING DATE: 2000-09-20
PRIOR APPLICATION NUMBER: 60/156,277
PRIOR FILING DATE: 1999-09-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: FastSEQ for Windows Version 4.0
EQ ID NO 10
LENGTH: 18
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TITLE OF INVENTION: SECRETED FACTORS
FILE REFERENCE: SCIOS.013A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Stanton, APPLICANT: Kapoun,
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PRIOR APPLICATION NUMBER: US/09/637,751
PRIOR FILING DATE: 2000-08-11
NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS:
                PRIOR APPLICATION NUMBER: 60/136,545
PRIOR FILING DATE: 1999-05-28
NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
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ORGANISM: Artificial Sequence
FEATURE:
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nes 18; Conservative (
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Ann Marie
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100.0%; Pred. No.
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100.0%; Pred. No. 5.4
ive 0; Mismatches
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. 5.4e+02;
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GENERAL INFORMATION:
APPLICANT: Brown, Sherri M.
APPLICANT: Heck, Gregory R.
APPLICANT: Piller, Kenneth J.
APPLICANT: Kishore, Ganesh M.
APPLICANT: Elich, Tedd D.

US-09-371-307-85/c

Sequence 85, Application US/09371307A Patent No. US20020053095A1

RESULT 62

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Best Local S
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Best Local Simi
Matches 18;
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                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 2, Application US/09917138 Patent No. US20020031776A1
                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/917,138
CURRENT FILING DATE: 2001-07-26
PRIOR APPLICATION NUMBER: 09/580,358
PRIOR FILING DATE: 2000-05-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: TULLIS, Richard
APPLICANT: STEIFFEL, Jerome
TITLE OF INVENTION: ENZYMATIC LABELIING AND DETECTION OF DNA
TITLE OF INVENTION: HYBRIDIZATION PROBES
FILE REFERENCE: 24730-2207B
                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/136,545
PRIOR FILING DATE: 1999-05-28
NUMBER OF SEQ ID NOS: 7
SOFTWARE: FASTSEQ for Windows Version 4.0
                                                                                                                                                                                                                           LENGTH: 19
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
                                                                                                                                                       OTHER INFORMATION: Oligonucleotide Primer NAME/KEY: misc_feature LOCATION:
OTHER INFORMATION: Combined DNA/RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LOCATION:
OTHER INFORMATION: Combined DNA/RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: modified_base
LOCATION: (1)
OTHER INFORMATION: Biotinylation at the 5'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: Oligonucleotide Primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence
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                                 2662 AAAAAAAAAAAAAAAAA 2679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2662 АЛАЛАЛАЛАЛАЛАЛА 2679
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1 AAAAAAAAAAAAAAAA 18
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18; Conservative (
                                                                                    Similarity
                                                                0.7%; Score 18; DB 9; Larity 100.0%; Pred. No. 5.4e+02; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.7%; Score 18;
100.0%; Pred. No.
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                                                                                                     Length 19;
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                                                                    Gaps
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RESULT 64
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CURRENT APPLICATION NUMBER: US/09/901,484A

CURRENT FILING DATE: 2001-07-09

PRIOR APPLICATION NUMBER: US 08/96,306

PRIOR FILING DATE: 1997-12-22

PRIOR APPLICATION NUMBER: US 60/099,658

PRIOR FILING DATE: 1998-09-09

PRIOR APPLICATION NUMBER: US 09/218,207

PRIOR APPLICATION NUMBER: US 09/218,207

PRIOR FILING DATE: 1998-12-22

PRIOR FILING DATE: 1999-06-23

PRIOR FILING DATE: 1999-06-23

PRIOR FILING DATE: 1999-06-23

PRIOR FILING DATE: 1999-06-23

PRIOR FILING DATE: 2001-05-11
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                                                                                                                                                                                                                                                                                             NUMBER PATE
SOFTWARE: Pate
; SEQ ID NO 515
FURTH: 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: Description of Artificial Sequence:Primer US-09-371-307-85
                                                                                                                 Matches
                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/371,307A
CURRENT FILING DATE: 1999-08-10
NUMBER OF SEQ ID NOS: 89
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 85
LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
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                                                                                                                                                                                                                                                                                 LENGTH: 19
                                                                                                                                                                              FEATURE:
NAME/KEY: misc feature
LOCATION: (1). (19)
OTHER INFORMATION: potential microsequencing oligo for 4-4-187.mis2
                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Bougueleret, Lydie
TITLE OF INVENTION: Prostate Cancer Gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Conen,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Rao, Sudabathula
                                                                                                                                                                                                                                                              ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: MOBT:216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPLICANT: Logusch, Sherry J.
FITLE OF INVENTION: Methods for controlling gibberellin levels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
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                                                                                                                                 Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                            2662 АДАДАДАДАДАДАДАД 2679
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18
                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                  PatentIn version 3.1
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Chumakov, Ilya
                                                                                                                 Conservative
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                                                                                                                             0.7%;
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                                                                                                                             Score 18;
Pred. No.
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Query Match
Best Local Similarity
Watches 18; Conserve
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TITLE OF INVENTION: PROSTATE CANCER GENE
FILE REFERENCE: GENSET.18CP1CP
CURRENT APPLICATION NUMBER: US/09/853,526
CURRENT FILING DATE: 2001-05-11
PRIOR APPLICATION NUMBER: 09/338,907
PRIOR FILING DATE: 1999-06-23
PRIOR PPLICATION NUMBER: 08/996,306
PRIOR APPLICATION NUMBER: 08/996,306
PRIOR FILING DATE: 1997-12-22
PRIOR PILING DATE: 1997-12-22
PRIOR APPLICATION NUMBER: 60/099,658
PRIOR FILING DATE: 1998-09-09
PRIOR APPLICATION NUMBER: 09/218,207
PRIOR APPLICATION NUMBER: 09/218,207
PRIOR APPLICATION NUMBER: 09/218,207
PRIOR FILING DATE: 1998-12-22
                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 34
SOFTWARE: Patentin version 3.1
SEQ ID NO 15
LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 15, Application US/09970971A Publication No. US20030096979A1 GENERAL INFORMATION:
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LENGTH: 19
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ORGANISM: Homo Sapiens
FEATURE:
NAME/KEY: misc feature
LOCATION: 1..19
                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/970,971A CURRENT FILING DATE: 2002-05-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Kawasaki, Andrew M.
TITLE OF INVENTION: Oligonucleotides Having
FILE REFERENCE: ISIS4789
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
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                                                                                                                                                                                      NAME/KEY: misc feature
LOCATION: (16)...(19)
OTHER INFORMATION: 3'-O-MOE
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                                                                                                                LOCATION: (1). (19)
OTHER INFORMATION:
                                                                                                                                  NAME/KEY: misc feature LOCATION: (1)...(19)
                                                                                                                                                                                                                                                                                                                              LENGTH: 19
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                                                                                                                                                                            FEATURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: potential microsequencing oligo for 4-4-187.mis2
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                                                                                                                                                                                                                                                                    OTHER INFORMATION: No.
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Cook, Phillip Dan
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Ilya, Chumakov
              0.7%; Score 18; DB 10; 1 ilarity 100.0%; Pred. No. 5.4e+02; Conservative 0; Mismatches 0;
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100.0%; Pred. No.
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5.4e+02;
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DB 10;

Length 19;

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APPLICANT: Cook, Phillip Dan
APPLICANT: Kawasaki, Andrew M.
TITLE OF INVENTION: Oligonucleotides Having Di
FILE REFERENCE: ISIS4789
CURRENT APPLICATION NUMBER: US/09/970,971A
CURRENT FILING DATE: 2002-05-03
NUMBER OF SEQ ID NOS: 34
SOPTWARE: Patentin version 3.1
SEQ ID NOS: 34
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US-09-970-971A-16/c
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US-09-970-971A-26
                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/970,971A
CURRENT FILING DATE: 2002-05-03
NUMBER OF SEQ ID NOS: 34
SOFTWARE: PatentIn version 3.1
SEQ ID NO 26
LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                        Sequence 26, Application US/09970971A Publication No. US20030096979A1
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                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                      APPLICANT: Manoharan, Muthiah
APPLICANT: Mohan, Venkatramar
APPLICANT: Cook, Phillip Dan
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        NAME/KEY: misc_feature
LOCATION: (16)...(19)
OTHER INFORMATION: 2'-modified T
                                                               FEATURE:
OTHER INFORMATION: NO.
FEATURE:
                                                                                                               TYPE: DNA
ORGANISM: Artificial Sequence
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OTHER INFORMATION: P=O
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LOCATION: (16)..(19)
OTHER INFORMATION: 3'-O-MOE
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OTHER INFORMATION: No.
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Similarity 100.0%; Pred. No.
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Cook, Phillip Dan
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                                                                                 US20030096979Alel Sequence
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; FEATURE:
, NAME/KEY: misc_feature
; LOCATION: (19)...(19)
; OTHER INFORMATION: N= G-clamp modification
US-09-996-292A-55
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; NAME/KEY: misc feature
; LOCATION: (19)...(19)
; OTHER INFORMATION: N= phenoxazine
US-09-996-292A-54
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                                                                                                        NUMBER OF SEQ ID NOS: 55
SOFTWARE: Patentin version 3.1
SEQ ID NO 55
LENGTH: 19
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Completely synthetic sequence
                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Mancharan, Muthiah
APPLICANT: Mancharan, Thazha P.
APPLICANT: Prakash, Thazha P.
APPLICANT: Rajeev, Kallanthottathil Gopalan
TITLE OF INVENTION: Nuclease Resistant Chimeric Oligonucleotides
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Publication No.
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Best Local Similarity
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APPLICANT: Rajeev, Kallanthottathil Gopalan
TITLE OF INVENTION: Nuclease Resistant Chimeric Oligonucleotides
FILE REFERENCE: ISIS-4804
CURRENT APPLICATION NUMBER: US/09/996,292A
CURRENT FILING DATE: 2001-09-28
NUMBER OF SEQ ID NOS: 55
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                         FILE REFERENCE: ISIS-4804
CURRENT APPLICATION NUMBER: US/09/996,292A
CURRENT FILING DATE: 2001-09-28
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APPLICANT: Maier, Martin A.
APPLICANT: Prakash, Thazha P.
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OTHER INFORMATION: Completely synthetic sequence
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ORGANISM: Artificial Sequence
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No. US20030158403A1
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100.0%; Pred. No.
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Query Match

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Length 19;

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Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 26
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 25
LENGTH: 19
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Publication No. US20020182687A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                 APPLICANT: Kurz, Markus
APPLICANT: Lohse, Peter
APPLICANT: Wagner, Richard
TITLE OF INVENTION: Peptide Acceptor Ligation Methods
FILE REFERENCE: 50036/031002
CURRENT APPLICATION NUMBER: US/10/208,357
CURRENT FILING DATE: 2002-07-30
PRIOR APPLICATION NUMBER: US/09/619,103
PRIOR APPLICATION NUMBER: US/09/619,103
PRIOR FILING DATE: 2000-07-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 3, Application US/10096221 Publication No. US20020164628A1 GENERAL INFORMATION:
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                                                                                              OTHER INFORMATION: designed sequence for nucleic acid purification .10-208-357-25
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                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/145,834 PRIOR FILING DATE: 1999-07-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PILE REFERENCE: 492692000700

CURRENT APPLICATION NUMBER: US/10/096,221

CURRENT FILING DATE: 2002-06-27

PRIOR APPLICATION NUMBER: US 60/274,236

PRIOR FILING DATE: 2001-03-09

NUMBER OF FILING DATE: 2001-03-09
                                                                                                                                 TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Kurn, Nurith
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR
TITLE OF INVENTION: AMPLIFICATION OF RNA SEQUENCES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS:
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ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Primer
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NAME/KEY: misc_feature
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                Conservative
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        0.7%; Score 18; DB 13; I
100.0%; Pred. No. 5.4e+02;
... Mismatches 0;
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. 5.4e+02;
                                               Length 19;
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          Indels
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US-10-123-597-2
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                                                                                                                                                        SOFTWARE: PatentIn version 3.1
SEQ ID NO 2
LENGTH: 19
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APPLICANT: Cook, Phillip D
APPLICANT: Kawasaki, Andre
APPLICANT: Manoharan, Muth
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NUMBER OF SEQ ID NOS: 28
SOFTWARE: PatentIn version 3.1
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 2, Application US/10123597
Publication No. US20030078415A1
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Best Local Similarity
                                                                                                                                                                                                                         FILE REFERENCE: ISIS5040
CURRENT APPLICATION NUMBER: US/10/123,597
CURRENT FILING DATE: 2002-07-10
PRIOR APPLICATION NUMBER: 09/227,782
PRIOR FILING DATE: 1999-01-08
NUMBER OF SEQ ID NOS: 28
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Publication No. US20030078415A1
                                                                                                                                                                                                                                                                                                                                        APPLICANT: Manoharan, Muthiah
APPLICANT: Prakash, Thazha P
APPLICANT: Fraser, Allister S
TITLE OF INVENTION: Regioselective Synthesis of 2'-0-Modified Nucleosides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/123,597
CURRENT FILING DATE: 2002-07-10
PRIOR APPLICATION NUMBER: 09/227,782
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APPLICANT: Fraser, Allister S
TITLE OF INVENTION: Regioselective Synthesis of 2'-O-Modified Nucleosides
FILE REFERENCE: ISIS5040
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APPLICANT: Kawasaki, Andre
APPLICANT: Manoharan, Mutl
                                    NAME/KEY: misc feature LOCATION: (15)..(18)
             OTHER INFORMATION: 5-methyl-2'dimethylaminooxyethoxy
                                                                       OTHER INFORMATION: Description of Artificial Sequence:
                                                                                                                                           TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic construct NAME/KEY: misc_feature
LOCATION: (15)..(18)
OTHER INFORMATION: 5-methyl-2'-aminooxyethoxy
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                                                                                                                ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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Manoharan, Muthiah
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Prakash, Thazha P
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US-10-123-597-4/c
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US-10-123-597-3/c
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APPLICANT: Mancharan, Muthiah
APPLICANT: Prakash, Thazha P
APPLICANT: Prakash, Thazha P
APPLICANT: Prakash, Thazha P
APPLICANT: Prakash, Thazha P
APPLICANT: SISSO40
CURENT APPLICATION: Regioselective Synthesis of 2'-O-Modified Nucleosides
FILE REFERENCE: ISISSO40
CURRENT APPLICATION NUMBER: US/10/123,597
CURRENT APPLICATION NUMBER: 09/27,782
PRIOR APPLICATION NUMBER: 09/227,782
PRIOR APPLICATION NUMBER: 09/227,782
PRIOR FILING DATE: 1999-01-08
NUMBER OF SEQ ID NOS: 28
SOFTWARE: Patentin version 3.1
SQ ID NO 4
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SEQ ID NO 3

SEQ ID NO 3

SEQ ID NO 3

SEQ ID NO 6

TYPE: DNA

ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic construct
NAME/KEY: misc feature
LOCATION: (15)...(18)
OTHER INFORMATION: 2'-methoxyethoxy
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Best Local Similarity
Matches 18; Conserv
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APPLICANT: Manoharan, Muthiah
APPLICANT: Prakash, Thazha P
APPLICANT: Fraser, Allister S
TITLE OF INVENTION: Regioselective Synthesis of 2'-O-Modified Nucleosides
FILE REFERENCE: ISISS040
CURRENT APPLICATION NUMBER: US/10/123,597
CURRENT FILING DATE: 2002-07-10
PRIOR APPLICATION NUMBER: 09/227,782
PRIOR FILING DATE: 1999-01-08
NUMBER OF SEQ ID NOS: 28
CONTRADED TO THE CONTRADED TO 
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Best Local (
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APPLICANT: Kawasaki, Andre
APPLICANT: Manoharan, Muth
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APPLICANT: Kawasaki, Andre
APPLICANT: Manoharan, Muth
OTHER INFORMATION: Description of Artificial Sequence: Synthetic construct NAME/KEY: misc_feature LOCATION: (16)...(19)
                                                                                                         TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                              ENGTH: 19
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. 5.4e+02;
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; OTHER INFORMATION: 5-methyl-2'-dimethylaminooxyethoxy US-10-123-597-4
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Query Match 0.7%; Score 18; DB 14; Length 19; Best Local Similarity 100.0%; Pred. No. 5.4e+02; Matches 18; Conservative 0; Mismatches 0; Indels 0;

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Search completed: February 3, 2005, 00:11:19 Job time: 1480.59 secs

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Title:
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/cgn2_6/ptcdata/1/ina/6B_COMB.seq:*
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PCT-US94-05407-7
PCT-US94-05407-10
US-09-318-907-10
US-09-218-207-10
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US-09-803-263-7
US-09-947-311-6
US-09-947-962-27
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Sequence 42, Application US/08113646A Patent No. 5578468 GENERAL INFORMATION:
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APPLICANT: PATEL, Dhavalkumar
APPLICANT: ANTCZAK, James B.
TITLE OF INVENTION: SITE-SPECIFIC RNA CLEAVAGE
NUMBER OF SEQUENCES: 44
CORRESSPONDENCE ADDRESS:
                COMPUTER READABLE
                                                                                                                ADDRESSEE: NIXON & VANDERHYE P.C. STREET: 1100 NORTH GLEBE ROAD CITY: ARLINGTON
                                         COUNTRY: U.S.A. ZIP: 22201-4714
                                                                                             VIRGINIA
                FORM:
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US-09-227-782-25
US-09-288-679-1
US-09-612-531-3
US-09-612-531-3
US-09-612-531-3
US-09-612-531-3
US-09-612-531-3
US-09-142-212A-10
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US-09-349-040A-3
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US-09-349-040A-5
US-10-123-597-1
US-10-123-597-2
US-10-123-597-3
US-10-123-597-1
US-09-349-033A-1
US-09-349-033A-1
US-09-371-373-593-5
US-09-371-373-593-5
US-08-937-080-83
US-08-937-080-83
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US-08-937-080-83
US-08-937-080-83
US-08-937-362-83
US-08-937-362-83
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COUNTEX: USA.
ZIP: 94306-2155
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve:
CURRENT APPLICATION NUMBER: US/08/455,627
FILING DATE: 31-MAY-1995
CLASSIFICATION: 435
CLASSIFICATION: 435
NAME: Nakamura, Jackie N.
REGISTRATION UMBER: 35,966
REFERENCE/DOCKET NUMBER: LYNX-003/(
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION ON BER 141-843-500
TELEFAX: 415-857-0663
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 30 nuclecides
TYPE: nucleic acid
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US-08-455-627-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No. 5571677

GENERAL INFORMATION:
APPLICANT: Sergei M. Gryaznov
TITLE OF INVENTION: Convergent Synthesis of Branched and Multiply
TITLE OF INVENTION: Connected Macromolecular Structures
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
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TELEPAN: (703) 816-4100
TELEPAN: (703) 816-4100
TELEY X: (703) 816-4100
TELEY X: 200797 NIXN UR
INFORMATION FOR SEQ ID NO: 42:
SEQUENCE CHARACTERISTICS:
LENGTH: 25 base pairs
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ver
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/113,646A
FILLING DATE: 31-AUG-1993
CLASSIFICATION DATA:
DATE: US 07/084,406
FILING DATE: 10-AUG-1987
ATTORNEY/AGENT INFORMATION:
NAME: WILSON, MARY J.
REGISTRATION NUMBER: 32,955
REGISTRATION NUMBER: 32,955
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
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CITY: Palo Alto
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; TOPOLOGY: linear
; MOLECULE TYPE: DNA
US-08-455-627-12
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Best Local Similarity 100.0%; P
Matches 20; Conservative 0;
                           Sequence 12, Application US/08787321A Patent No. 6180777
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Best Local
GENERAL INFORMATION:
APPLICANT: Horn, Thomas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patent No.
                                                                                                                                                                                                                                                                                                                                           TELEFAX: 415-857-0663
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08
FILING DATE: 31-MAY-1995
ATTORNEY/AGENT INFORMATION:
NAME: Nakamura, Jackie N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/689,856
                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 30 nucleotides
TYPE: nucleic acid
                                                                                                                                                                                                                                                      MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: 415-843-5000
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TOPOLOGY: 111
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REFERENCE/DOCKET NUMBER: LYNX-003/01 US
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
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Five Palo Alto Square,
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L00.0%; Pred. No.
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                                                                 US-08-482-918-32
   Query Match 0.7%; Sco
Best Local Similarity 100.0%; P
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Best Local Similarity
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LENGTH: 30
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                                                                                                                                                        TELEPHONE: 312/474-630
TELEPAX: 312/474-0448
TELEX: 25-3856
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/0
FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: 312/474-6300
                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Suggs, Sidney V. APPLICANT: Martin, Francis H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                                                 MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION:
                                                                                                                                                                                                                                                     NAME: Clough, David W. REGISTRATION NUMBER: 36 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: United States of America
                                                                                              TOPOLOGY:
                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 424
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                                                                                                                                         LENGTH:
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                                                                                                                         nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                          READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3: Marshall, O'Toole, 6300 Sears Tower, 233
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Suggs, Sidney V.
                                                                                                                                                                                                                                                                                                                                                                                          E: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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100.0%; Pred. No.
tive 0; Mismatc
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                                                                                                                                                                                                                                                                 36,107
    ; Score 19; DB
; Pred. No. 2.3
0; Mismatches
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DB 3; Lt...
5. 2.3e+02;
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US-09-224-681-32
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US-09-224-681-32/c
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Patent No.
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FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/005,893
FILING DATE: 12-JAN-1998
CLASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                       INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                              NAME: Clough, David W.
REGISTRATION NUMBER: 36,107
REFERENCE/DOCKET NUMBER: 010
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312/474-6300
              TOPOLOGY: li
                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: 07/4:
FILING DATE: 16-OCT-1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 07/537,198
FILING DATE: 11-JUN-1990
PRIOR APPLICATION DATA:
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APPLICATION NUMBER: 07/982,255
FILING DATE: 25-NOV-1992
PRIOR APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 07/573,616
FILING DATE: 24-AUG-1990
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: 07/589,701 FILING DATE: 01-OCT-1990 PRIOR APPLICATION DATA:
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ADDRESSEE: Marshall, O'Toole,
STREET: 6300 Sears Tower, 233
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APPLICANT: Suggs, Sidney V.
APPLICANT: Martin, Francis H.
TITLE OF INVENTION: Method for Enhancing the Efficiency of Gene
TITLE OF INVENTION: Transfer with Stem Cell Factor (SCF) Polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                  STRANDEDNESS:
                                                                                                                                                                           TELEFAX: 312/474-0448
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FILING DATE: 24-MAY-1995
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                                                                       nucleic acid
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                                                                                           20 base pairs
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                                      linear
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South Wacker Drive
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; TOPOLOGY: 11; MOLECULE TYPE: US-08-336-728A-32
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Query Match
Best Local Similarity 100.0%; P
Matches 19; Conservative 0;
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Best Local S
Matches 19
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                 PILING DATE: 01-OCT-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/573
FILING DATE: 24-AUG-1990
PRIOR APPLICATION UNMBER: 07/537
FILING DATE: 11-JUN-1990
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 07/432
FILING DATE: 11-GORNATION:
APPLICATION UNMBER: 07/422
FILING DATE: 16-OCT-1989
ATTORNEY/AGENT INFORMATION:
NAME: Clough, David W.
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                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 01
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312/474-6300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING CONTROL OF THE SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
CURRENT APPLICATION NUMBER: US/08/336,728A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 07/91
FILING DATE: 25-NOV-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/51
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FITLE OF INVENTION: Stem Cell Factor
                                                                                                                           TYPE: nucleic acid
TYPE: single
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                         Clough, David W.
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Bosselman, Robert A.
Suggs, Sidney V.
Martin, Francis H.
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               0.7%; Score 19; DB 3;
100.0%; Pred. No. 2.3e+0
ive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                 07/422,383
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                             2.3e+02;
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                                               Length 20;
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RESULT 8
US-09-635-251-32/c
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                                                                                                                                   SEQUENCE DESCRIPTION: SEQ ID NO: US-09-635-251-32
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                                                           Matches
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                                                                         Local
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APPLICATION NUMBER: US/09/635,251
FILING DATE: 07-Aug-2000
                                                                                                                                                                                                                                                                                                                                                              NAME: Clough, David W.
REGISTRATION NUMBER: 36,107
REFERENCE/DOCKET NUMBER: 011
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/449,182
FILING DATE: 24-MAY-1995
APPLICATION NUMBER: 08/172,329
FILING DATE: 21-DEC-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2661
2661 CAAAAAAAAAAAAAAAA 2679
                                                                                                                                                                           TOPOLOGY: linear MOLECULE TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
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                                                           l Similarity
19; Conserv
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                             LENGTH: 20 base pairs TYPE: nucleic acid
                                                                                                                                                                                                             STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                            TELEPHONE: 312/474-6300
TELEFAX: 312/474-0448
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 07/422,383 FILING DATE: 16-OCT-1989
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                                                                                                                                                                                                                                                                                                            TELEX: <Unknown>
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                                                     0.7%; Score 19; DB 4; L
larity 100.0%; Pred. No. 2.3e+02;
Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CATION NUMBER: 07
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South Wacker Drive
                                                                                           Length 20;
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Best Local Similarity
APPLICATION NUMBER: 08/0
ETIJING DATE: 13-MAY-1993
INFORMATION FOR SEQ ID NO: 88
SEQUENCE CHARACTERISTICS:
LENGTH: 23 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 7: SEQUENCE CHARACTERISTICS:
                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
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CURRENT APPLICATION DATA:
APPLICATION UMBER: PCT/US94/05407
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                  APPLICATION NUMBER: I
                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 1. CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear MOLECULE TYPE: oligonucleotide
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                                                                                                                                                                                                                                                                         STREET: Suite
CITY: Atlanta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 08/0 FILING DATE: 13-MAY-1993
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Y: USA
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Suite 1200, 127 Peachtree Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                E: NEEDLE & ROSENBERG, P.C.
Suite 1200, 127 Peachtree Street
                                                                                                                                                                                                                                              USA
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100.0%; Pred. No.
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nucleic acid

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US-08-996-306-10/c
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Best Local Similarity
Marches 19; Conserve
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                                                                                Query Match
                                                      Matches
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                                                                                                                                                                                                                                                                                            TELEFAX: (619) 235-0176
INFORMATION FOR SEQ ID NO: 10:
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                                                                                                                  FEATURE:
NAME/KEY: PGRT32
LOCATION: complement 5198..5221
OTHER INFORMATION: Location rel
                                                                                                                                                                                              MOLECULE TYPE:
ORIGINAL SOURCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Cohen, Daniel
APPLICANT: Chumakov, Ilya
APPLICANT: Blumenfeld, Marta
APPLICANT: Bougheleret, Lydie
TITLE OF INVENTION: Prostate cancer gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: oligonucleotide
                                                                                                                                                                                                                                                                                                                                    NAME: Israelsen, Ned A.
REGISTRATION NUMBER: 29,655
REFERENCE/DOCKET NUMBER: GE
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/996,306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy Disk
                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
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                                                                  госат
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible OPERATING SYSTEM: Win95
                                                                                                                                                                                                                          TOPOLOGY:
                                                                                                                                                                                                                                    STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Word
                                                                                                                                                                                 DRGANISM:
                                                                                                                                                                                                                                                                                                                         TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Knobbe,
                                                                                                                                                                                                                                                                  ENGTH:
                       2661 САЛАЛАЛАЛАЛАЛАЛА 2679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2661 СААЛАЛАЛАЛАЛАЛАЛА 2679
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                                                                  Similarity
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San Diego
California
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                                                                                                                                                                                                                                                                  24 base pairs
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                                                    Conservative
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                                                                0.7%; Score 19;
100.0%; Pred. No.
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                                                                                                                      Location relative to seqID3
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100.0%; Pred. No. 2.:
ive 0; Mismatches
                                                   0;
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                                                   Mismatches
                                                 DB 2; Length 24;
2.3e+02;
hes 0; Indels
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RESULT 12

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2661 CAAAAAAAAAAAAAAAAA 2679

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US-09-218-207-10
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APPLICANT: Bougueleret, Lydie
TITLE OF INVENTION: Prostate cancer gene
FILE REFERENCE: GENSET.018CP1
CURRENT APPLICATION NUMBER: US/09/218,207
CURRENT FILING DATE: 1998-12-22
EARLIER APPLICATION NUMBER: 08/996,306
EARLIER APPLICATION NUMBER: 08/996,306
EARLIER APPLICATION NUMBER: 60/099,658
EARLIER APPLICATION NUMBER: 60/099,658
EARLIER FILING DATE: 1998-09-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-218-207-10/c
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CURRENT FILING DATE: 1999-06-23
EARLIER APPLICATION NUMBER: 08/996,306
EARLIER FILING DATE: 1997-12-22
EARLIER APPLICATION NUMBER: 60/099,658
EARLIER FILING DATE: 1998-09-09
EARLIER APPLICATION NUMBER: 69/218,207
EARLIER FILING DATE: 1998-12-22
Query Match
Best Local Similarity 100.
19; Conservative
                                                                                                                                                                                                             SEQ ID NO 10
LENGTH: 24
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LENGTH: 24
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                                                                                                                                                                                                                                              SOFTWARE: Patent.pm
                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Cohen, Daniel APPLICANT: Blumenfeld, M
                                                                                           NAME/KEY: misc_feature
LOCATION: 1..24
OTHER INFORMATION: primer oligonucleotide PGRT32
                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 578
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SOFTWARE: Patent.pm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Cohen, Daniel APPLICANT: Blumenfeld, M
                                                                                                                                                                         TYPE: DNA
ORGANISM: Homo Sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: PROSTATE CANCER GENE
FILE REFERENCE: GENSET.18CP1CP
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                                                                                                                                                              FEATURE:
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ORGANISM: Homo Sapiens
FEATURE:
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         0.7%; Urr
100.0%; Prr
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100.0%; Pred. No. 2.3
ive 0; Mismatches
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                                  Score 19;
Pred. No.
                  Mismatches
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o. 2.3e+02;
0;
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                Indels
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RESULT 15
US-08-621-914A-3/c
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US-09-721-154-6/c
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Patent No. 6651008
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 6
LENGTH: 24
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APPLICANT: Adams, Cynthia
APPLICANT: Sabry, James
APPLICANT: Crompton, Anne
                                                                                                                                                                                                                            ZIP: 10036-2711

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, V
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:

APPLICANT: KATO, KIKUYA
TITLE OF INVENTION: MOLECULAR INDEXING FOR EXPRESSED GENE
TITLE OF INVENTION: ANALYSIS
NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
ADDRESSEE: PENNIE & EDMONDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Database system including computer code TITLE OF INVENTION: for predictive cellular bioinformatics FILE REFERENCE: Cytop007C2 CURRENT APPLICATION NUMBER: US/09/721,154 CURRENT FILING DATE: 2002-06-14 PRIOR APPLICATION NUMBER: 09/311,996 PRIOR APPLICATION NUMBER: 09/311,996 PRIOR APPLICATION NUMBER: 09/311,996 PRIOR FILING DATE: 1999-05-14 NUMBER OF SEQ ID NOS: 14
                                                                      APPLICATION NUMBER: US/08/621,91
FILING DATE: 26 MAR-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: LAWRENCE III, STANTON T.
REGISTRATION NUMBER: 25,736
REFERENCE/DOCKET NUMBER: 7005-10
TELECOMMUNICATION INFORMATION:
TELEX: 66141 PENNIE INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: Psuedo-sequence
                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 1155 A
                                    TELEFAX:
                                                      TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2660 АСАЛАЛАЛАЛАЛАЛАЛА 2678
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                                    : (212) 790-90
(212) 869-9741
                                                                                                                                                                                                                                                                                                                                                                    USA
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100.0%; Pred. No.
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                                                                                              7005-107-999
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. 2.3e+02;
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Sequence 7, Application US/10003998A
Patent NO. 6664064
GENERAL INFORMATION:
APPLICANT: Roche Diagnostics GmbH
TITLE OF INVENTION: Method for melting curve
TITLE OF INVENTION: products
FILE REFERENCE: 5438/00/EP
CURRENT APPLICATION INVESER: US/10/003,998A
CURRENT FILING DATE: 2001-11-14
NUMBER OF SEQ ID NOS: 8
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US-10-003-998A-7
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                                                                                                                                                                                                               APPLICANT: Fisher, Paul B.
APPLICANT: Kang, Dong-Chul
TITLE OF INVENTION: METHOD FOR FULL-LENGTH CDNA CLONING
TITLE OF INVENTION: USING DECEMERATE STEM LOOP ANNEALIN
FILE REFERENCE: A34701 (070050.1728)
CURRENT FILING DATE: US/09/724,028A
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 33
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 18
                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.7%; Score 19;
Best Local Similarity 100.0%; Pred. No.
Matches 19; Conservative 0; Mismatc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 7
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Best Local Similarity
Matches 19; Conserva
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                                                                  Query Match
Best Local
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TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                         TYPE: DNA
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                     OTHER INFORMATION: Oligonucleotide primer PCTAR3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: '26 base pairs
TYPE: nucleic acid
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2660 ACAAAAAAAAAAAAAAAA 2678
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                                                                    Similarity
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                                              0.7%; Score 19; DB 4; 100.0%; Pred. No. 2.3e+0 tive 0; Mismatches
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                                                                    2.3e+02;
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US-08-621-914A-16/c
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                                                                                                                                                                                                                                                              Sequence 3, Application US/08346429 Patent No. 5837820
                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                            Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
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                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: DeRose,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEX: 66141 PENNIE INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: KATO, KIKUYA
TITLE OF INVENTION: MOLECULAR
TITLE OF INVENTION: ANALYSIS
                                                                                      TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                            APPLICANT: Duval, Manuel APPLICANT: Job, Claudette APPLICANT: Job, Dominique
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: LAWRENCE II, STANTON T.
REGISTRATION NUMBER: 25,736
REFERENCE/DOCKET NUMBER: 7005-107-999
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEPHONE: (212) 790-9090
                                                       CORRESPONDENCE ADDRESS:
                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: unknown MOLECULE TYPE: other nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 10036-2711

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/621,914A
FILING DATE: 26-MAR-1996
                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESSE: PENNIE & EDMONDS
STREET: 1155 AVENUE OF THE AMERICAS
                     STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS:
                                    ADDRESSEE:
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o. 5707807
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SEE: SCULLY SCOTT MURPHY & PRESSER
1: 400 Garden City Plaza
Garden City
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                                                                                                                                                                                   DeRose, Richard
Douce, Roland
Duval, Manuel
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100.0%; Pred. No. 6.1
ative 0; Mismatches
                                                                                        PROTEIN CAPABLE OF BEING BIOTINYLATED WHICH CAN
BE USED FOR DETERMINING THE GERMINATION STAGE OF
A SEED
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hes 0;
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/346,429
FILING DATE: 29-NOV-1994
CCLASSIFICATION: 530
ATTORNEY/AGRAM
                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/358,556A
FILING DATE: 14-DEC-1994
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
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Best Local (
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            APPLICATION NUMBER: FR 931516
FILING DATE: 16-DEC-1993
ATTORNEY/AGENT INFORMATION:
NAME: Player, William E.
REGISTRATION NUMBER: 31,409
REFERENCE/DOCKET NUMBER: 1057
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Chatelain, Franco
APPLICANT: Kunarev, Viktor
TITLE OF INVENTION: Process
TITLE OF INVENTION: a Solid
TITLE OF INVENTION: Implemer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: 516-742-4343
TELEFAX: 516-742-4366
TELEX: 230 901 SANS UR
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: DiGiglio, Frank S. REGISTRATION NUMBER: 31,346 REFERENCE/DOCKET NUMBER: 95 TELECOMMUNICATION INFORMATION: 516-742-4343
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                         CITY: WA
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STRANDEDNESS:
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DEDNESS: double
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O Seventh St. N.W.
(202) 638-6666
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                           Process for Preparing Polynucleotides on a Solid Support and Apparatus Permitting Implementation
                                                                                                                      FR 9315164
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TELEX: RCA 248593 IDEA INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:

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GENERAL INFURENCE Chatelain, APPLICANT: Chatelain, APPLICANT: Kumarev, Viktor TITLE OF INVENTION: A Solid TITLE OF INVENTION: A Solid TITLE OF INVENTION: Implement TITLE OF SEQUENCES: 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 21
US-08-358-556A-18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No. 586yow.
                                                                                 TELEFAX: (202) 393-5350
TELEX: RCA 248593 IDEA UR
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/358,556A
FILING DATE: 14-DEC-1994
CLASSIFICATION: 536
PRICR APPLICATION: 536
PRICR APPLICATION: 536
PRICR APPLICATION: 536
PRICR APPLICATION: 536
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                       FILING DATE: 16-DEC-1993
ATTORNEY/AGENT INFORMATION:
NAME: Player, William E.
REGISTRATION NUMBER: 31,409
REFERENCE/DOCKET NUMBER: 10577/P58418
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)638-6666
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: N-terminal
                                                TOPOLOGY: 11
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              HYPOTHETICAL:
ANTI-SENSE: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
les 18; Conserv
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LOCATION: 1...
                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Washington D.C
TYPE:
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                                                DNA (genomic)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Process for Preparing Polymucleotides on
a Solid Support and Apparatus Permitting its
Implementation
                                                                                                                                                                                                                                                                                                                                       FR 9315164
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RESULT 23 US-08-295-509B-4/c

Sequence 4, Application US/08295509B Patent No. 6045995

GENERAL INFORMATION:

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US-08-469-852A-4/c
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                                                                                                                                     US-08-469-852A-4
                                                               Matches 18; Conservative
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                                                                                 Best Local Similarity
                                                                                               Query Match
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                                                                                                                                                                                                                      TELEFAX: 215-568-3439
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/469,852A FILING DATE: 06-7UN-1995 CLASSIFICATION: 435 PRIOR APPLICATION DATA: 08/295,509 APPLICATION NUMBER: 08/295,509
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch disk, 1.44 Mb
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz & No. 5874213ris LLP STREET: One Liberty Place - 46th Floor
                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: IS
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-568-3100
                                                                                                                                                                                                                                                                                                                         FILING DATE: 24-AUG-1994
ATTORNEY/AGENT INFORMATION:
NAME: Michael P. Straher
REGISTRATION NUMBER: 38,3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: WordPerfect 6.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Capillary Electrophoretic TITLE OF INVENTION: Nucleic Acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Cummins, Lendell
APPLICANT: Freier, Susan M.
APPLICANT: Griffey, Richard
APPLICANT: Srivatsa, Susan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY:
                                                                                                                                                     TOPOLOGY:
                                                                                                                                                                                                      LENGTH:
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18 AAAAAAAAAAAAAAA 1
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                                                                                                                                                       linear
                                                                                 0.7%; Score 18;
100.0%; Pred. No.
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                                                                0; Mismatches
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                                                                                  6.1e+02;
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                                                                Indels
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US-08-884-029-9/c
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Best Local
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Michael P. Straher
REGISTRATION NUMBER: 38,325
REFERENCE/DOCKET NUMBER: ISIS
TELECOMMUNICATION INFORMATION:
TELECHONE: 215.568-3100
       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                NUMBER OF SEQUENCES: 9

CORRESPONDENCE ADDRESS:

ADDRESSEE: Townsend and Townsend and Crew LLP

STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                        APPLICANT: Lin, Ching-I Patsy
APPLICANT: Wallace, Robert Bruce
APPLICANT: Cossman, Jeffrey
APPLICANT: French, Cynthia
TITLE OF INVENTION: Lyophilization of Cultured Human Cells
TITLE OF INVENTION: to Preserve RNA and DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch disk, 1.44 Mb
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPerfect 6.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 24-AUG-19
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Srivatsa, Susan G.
TITLE OF INVENTION: Capillary Electrophoretic Detection
TITLE OF INVENTION: Nucleic Acids
APPLICATION NUMBER:
                                                                                                                                                COUNTRY:
                                                                                                                                                                    CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: One Liberty CITY: Philadelphia
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Similarity 100.0%;
18; Conservative (
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One Liberty Place - 46th Floor
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Srivatsa, Susan (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ss: single
linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-AUG-1994
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US/08/884,029
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Pred. No.
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RESULT 25
US-08-941-445A-30
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                                                                                                                                                                                                                                    OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, Ver CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/941,445A FILING DATE: 30-SEP-1997 CLASSIFICATIN: 800
PRIOR APPLICATION: 800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 30,
                         TELEPAX: (303) 499-808:
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
                                                                                                           PRIOR APPLICATION DATA:
APPLICATION UMMBER: US 60/026,855
FILING DATE: 30-SEP-1996
ATTORNEY/AGENT INFORMATION:
NAME: Winner, Ellen P
REGISTRATION UMBER: 28,547
REFERENCE/DOCKET NUMBER: 89-97
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 50 CITY: Boulder CO TIS
                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Keeling, Peter
APPLICANT: Guan, Hanping
TITLE OF INVENTION: Starch Encapsulation
NUMBER OF SEQUENCES: 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 02
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: DNA
                                                                                                  TELEPHONE:
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STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION:
OTHER INFORMATION:
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STRANDEDNESS: single
TOPOLOGY: linear
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100.0%; Pred. No. 6.1e+02;
tive 0; Mismatches 0;
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/note= "t at positions 13-18 may be
present or absent"
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RESULT 27
US-09-545-225-9/c
US-09-545-225-9/c
; Sequence 9, Application u...
; Patent No. 6410321
; GENERAL INFORMATION: Ching-I Patsy
Wallace, Robert Bruce
COSSMan, Jeffrey
"rench, Cynthia
"rench, Cynthia
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; MOLECULE TYPE:
; HYPOTHETICAL: N
US-08-941-445A-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 6
LENGTH: 18
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Feng, Li
APPLICANT: Latimer, Darin R.
TITLE OF INVENTION: Binary Encoded Sequence Tags
Patent No. 6383754
FILE REFERENCE: AGL 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
APPLICANT:
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CURRENT FILING DATE: 2000-08-11
NUMBER OF SEQ ID NOS: 10
SOPTWARE: Patentin Ver. 2.1
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les 18; Conservative 0;
ZIP: 94111-3834

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: JBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOPTWARE: PATENTIN Release #1.0,

CURRENT APPLICATION DATA:
                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                  TITLE OF INVENTION: Lyophilization of Cultured Human to Preserve RNA and DNA
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                                                                                                                                CITY: San Francisco
STATE: California
                                                                                                                     COUNTRY: USA
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100.0%; Pred. No. 6.1
ive 0; Mismatches
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100.0%; Pred. No. 6.1
ive 0; Mismatches
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hes 0;
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                   Version #1.30
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US-09-619-103-24
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                                                                                                                                                      US-09-619-103-24
                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 26
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 24
LENGTH: 18
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 24, Apprase No. 6429300
                                                                                              Query Match
Best Local 9
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Best Local Similarity 100.0%; Pred. No. 6.1e+02;
Matches 18; Conservative 0; Mismatches 0;
                                                                        Matches
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APPLICANT: Lohse, Peter
APPLICANT: Lohse, Peter
APPLICANT: Wagner, Richard
TITLE OF INVENTION: Peptide Acceptor Ligation Methods
FILE REFERENCE: 50036/031002
CURRENT APPLICATION NUMBER: US/09/619,103
CURRENT FILING DATE: 2000-07-19
PRIOR APPLICATION NUMBER: 60/145,834
PRIOR PILING DATE: 1999-07-27
                                                                                                                                                                  OTHER INFORMATION: designed sequence for nucleic acid purification
                                                                                                                                                                                           FEATURE:
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LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: modified_base
LOCATION: 13..18
OTHER INFORMATION: /mod_base= OTHER
/note= "t at positions 13-18 may be
present or absent"
SEQUENCE DESCRIPTION: SEQ ID NO: 9:
                    2662 AAAAAAAAAAAAAAAAA 2679
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APPLICATION NUMBER: US 08/884,029
FILING DATE: 27-JUN-1997
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                                                        Similarity 10.0%; Pred. No. ...
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FILING DATE: 07-Apr-2000
CLASSIFICATION: <Unknown>
ААААААААААААААА 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Application US/09619103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           576-0300
                                                                                          Score 18; DB 4; Li; Pred. No. 6.1e+02;
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                                                                                                          Length 18;
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RESULT 29 US-09-370-541-14/c

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RESULT 30
US-10-125-295-9/c
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US-09-370-541-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 9, Application Patent No. 6686460 GENERAL INFORMATION:
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EARLIER FILLING DATE: 1999-06-25
NUMBER OF SEQ ID NOS: 21
SOFTWARE: Patentin Ver. 2.0
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CURRENT APPLICATION NUMBER: US/09/370,541
CURRENT FILING DATE: 1999-08-09
EARLIER APPLICATION NUMBER: 09/130,973
EARLIER FILING DATE: 1998-08-07
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APPLICANT: Cook, Phillip Dan
APPLICANT: Frakash, Thatha P
APPLICANT: Kawasaki, Andrew M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EARLIER APPLICATION NUMBER: 09/016,520
EARLIER FILING DATE: 1998-01-30
EARLIER APPLICATION NUMBER: 60/037,143
EARLIER FILING DATE: 1997-02-14
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TITLE OF INVENTION: Compounds Prepared Therefrom
TITLE OF INVENTION: Compounds Prepared Therefrom
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
                                                                                CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/125,295
FILING DATE: 17-Apr-2002
CLASSIFICATION: -Unknown>
                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                   PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Lin, Ching-I Patsy
Wallace, Robert Bruce
Cossman, Jeffrey
French, Cynthia
                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Lyophilization of Cultured Human Cells to Preserve RNA and DNA
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APPLICATION NUMBER: US/09/545,225
FILING DATE: 07-Apr-2000
APPLICATION NUMBER: US 08/884,029
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                                                                                                                                                                                                                                                                                                                  STATE: California
                                                                                                                                                                                                                                                                                                   COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                        CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Application US/10125295
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Pred. No.
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                                                                                                                                                                     Version #1.30
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RESULT 32
US-09-803-263-7/c
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Sequence 7, Application US/09803263
Patent No. 6706476
GENERAL INFORMATION:
APPLICANT: Thirstrup, Kenneth
TITLE OF INVENTION: A Process for Amplifying and Labeling Single Stranded cDNA by
TITLE OF INVENTION: Adapter Mediated Amplification
FILE REFERENCE: 674513-2003.1
CURRENT APPLICATION NUMBER: US/09/803,263
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APPLICANT: Thirstrup, Kenneth
APPLICANT: Thirstrup, Kenneth
TITLE OF INVENTION: A Process for Amplifying and Labeling Single Stranded cDNA by
TITLE OF INVENTION: Adapter Mediated Amplification
FILE REFERENCE: 674513-2003.1
CURRENT APPLICATION NUMBER: US/09/803,263
CURRENT FILING DATE: 2001-03-09
NUMBER OF SEQ ID NOS: 19
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence PEATURE:
                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Poly-a tail
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                     Local Similarity
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SEQUENCE DESCRIPTION: SEQ ID NO: 9:
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TYPE: nucleic acid
STRANDEDNESS: single
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REGISTRATION NUMBER: 42,058
REFERENCE/DOCKET NUMBER: 02558B-059100US
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100.0%; Pred. No.
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100.0%; Pred. No.
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CURRENT FILING DATE:

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ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Complement of poly-a tail
US-09-803-263-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-994-311-6/c; Sequence 6, App.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-994-311-6
                                                                                                                                                                                                         Sequence 29, Application US/09142108C Patent No. 6774285 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Latimer, Darin R.
TITLE OF INVENTION: Binary Encoded Sequence Tags
Patent No. 6773886
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 19
SOFTWARE: PatentIn versionsEQ ID NO 7
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Best Local
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                       APPLICANT: Brugliera, Filippa
APPLICANT: Holton, Timothy A.
APPLICANT: Holton, Timothy A.
APPLICANT: Michael Michael Z.
TITLE OF INVENTION: GENETIC SEQUENCES ENCODING FLAVONOID PATHWAY ENZYMES
TITLE OF INVENTION: AND USES THEREFOR
FILE REFERENCE: 11658
CURRENT APPLICATION NUMBER: US/09/142,108C
CURRENT FILING DATE: 1998-09-01
PRIOR APPLICATION NUMBER: PN8386
PRIOR FILING DATE: 1996-03-01
NUMBER OF SEQ ID NOS: 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: AGL 100
CURRENT APPLICATION NUMBER: US/09/994,311
CURRENT FILING DATE: 2001-11-26
PRIOR APPLICATION NUMBER: US/09/637,751
PRIOR FILING DATE: 2000-08-11
NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No.
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SOFTWARE: P
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EQ ID NO 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 18
TYPE: DNA
ORGANISM: Artificial Sequence
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               PatentIn Ver. 2.1
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Lizardi, Paul M.
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Pred. No.
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Sequence 4, Application PC/TUS9405407
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; OTHER INFORMATION: Description
US-09-142-108C-29
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                                                                                                                                                                                                RESULT 36
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                                                                                                                                               Sequence 1, Application US/08756728A Patent No. 5821354
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Best Local
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Best Local S
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                                                                                                                                    GENERAL
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FILING DATE: 13-MAY-1993
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA ORGANISM: Artificial Sequence
                                                                   APPLICANT: Leclerc, Guy
APPLICANT: Martel, Remi
TITLE OF INVENTION: RAD
TITLE OF INVENTION: OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/05407
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Klauber & Jackson
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NUMBER OF SEQUENCES:
                                                       NUMBER OF SEQUENCES:
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OPERATING SYSTEM:
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COUNTRY: USA
            STREET:
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                                                                                                                                    INFORMATION:
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Hackensack
            411 Hackensack Avenue, 4th Floor
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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100.0%; Pred. No. 6.
tive 0; Mismatches
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100.0%; Pred. No.
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US-08-469-852A-2/c
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/756,728A
FILING DATE: 26-NOV-1996
CLASSIFICATION: 514
ATTORNEY/AGENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 2, Application US/08469852A Patent No. 5874213
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                                                                APPLICATION NUMBER: UPFILING DATE: 06-UN-11
CILASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 01
PRILITATION NUMBER: 01
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch disk, 1.44 Mb
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
             FILING DATE: 24-AUG-1994
ATTORNEY/AGENT INFORMATION:
NAME: Michael P. Straher
                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz & No. 5874213ris LLP
STREET: One Liberty Place - 46th Floor
                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Cummins, Lendell L.
APPLICANT: Freier, Susan M.
APPLICANT: Griffey, Richard
APPLICANT: Srivatsa, Susan G.
TITLE OF INVENTION: Capillary Electrophoretic Detection
TITLE OF INVENTION: Nucleic Acids
NUMBER OF SEQUENCES: 4
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HYPOTHETICAL:
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REGISTRATION NUMBER: 26,742
REFERENCE/DOCKET NUMBER: 1398-1-001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-487-5800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      / Match 0.7%; Score 18; DB Local Similarity 100.0%; Pred. No. 6. 18; Conservative 0; Mismatches
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                                                                                                                                PRIOR AFFILE.

APPLICATION NUMBER: 08/146,50*
FILING DATE: No. 6017696ember 1, 1993
ATTORNEY/AGENT INFORWATION:
NAME: Murphy, David B:
REGISTRATION NUMBER: 31.125
REFERENCE/DOCKET NUMBER: 207/263
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
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Best Local Similarity
Matches 18; Conserv
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                                                                                                      INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                     OPERATING SYSTEM: IBM P.C. DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/271,882B
FILING DATE: ULY 7, 1994
CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 655 ... CITY: LOS Angeles CTATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: SELF-ADDRESSABLE
TITLE OF INVENTION: SELF-ASSEMBLING
TITLE OF INVENTION: MICROSLECTRONIC SYSTEMS AND
TITLE OF INVENTION: MOLECULAR BIOLOGICAL ANALYSI
TITLE OF INVENTION: AND DIAGNOSTICS
NUMBER OF SEQUENCES: 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Michael
APPLICANT: Eugene 7
                                                                                                                     TELEPHONE: (213)
TELEFAX: (213)
TELEX: 67-3510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION: 215-568-3100
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STRANDEDNESS: sin
TOPOLOGY: linear
                                                      TYPE: nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: U
ZIP: 90071
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
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                                     acid
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633 West Fifth Street
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Glen A. Evans
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100.0%; Pro
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%; Pred. No. 6.1
0; Mismatches
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CORRESPONDENCE ADDRESS:

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US-08-881-784-18/c
Sequence 18, Application US/08881784; Patent No. 608731; GENERAL INFORMATION:
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Best Local Similarity
Matches 18; Conserv
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Best Local
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Patent No. 6045995
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch disk, 1.44 Mb
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Cummins, Lendell L.
APPLICANT: Freier, Susan M.
APPLICANT: Griffey, Richard
APPLICANT: Srivatsa, Susan G.
TITLE OF INVENTION: Capillary Electrophoratic Detection of
TITLE OF INVENTION: Nucleic Acids
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                APPLICANT: Croteau, Rodney B. APPLICANT: Lupien, Shari L. APPLICANT: Karp, Frank TITLE OF INVENTION: RECOMBINA TITLE OF INVENTION: THE PRODU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION: TELEPHONE: 215-568-3100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Michael P. Straher
REGISTRATION NUMBER: 38,325
REFERENCE/DOCKET NUMBER: ISIS-1395
                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: One Liberty
CITY: Philadelphia
STATE: PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/0 FILING DATE: 24-AUG-1994 CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH:
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RECOMBINANT MATERIALS AND METHODS FOR THE PRODUCTION OF LIMONENE HYDROXYLAS 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.7%; Score 18; DB 3; I
100.0%; Pred. No. 6.1e+02;
tive 0; Mismatches 0;
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                  HYDROXYLASES
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                                                                                         ; OTHER INFORMATION: Description of Artificial Sequence; OTHER INFORMATION: Sequence US-09-234-237-1
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Matches
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                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                               Matches
                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                    Sequence 1, Application US/09234237 Patent No. 6127124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                           APPLICANT: Leeds, Janet M
APPLICANT: Cummins, Lendell L
TITLE OF INVENTION: Fluorescence Based Nuclease Assay
FILE REFERENCE: ISIS3308
CURRENT APPLICATION NUMBER: US/09/234,237
CURRENT FILING DATE: 1999-01-20
RUMBER OF SEQ ID NOS: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (206) 224-07: INFORMATION FOR SEQ ID NO:
                                                                                                                                     TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                   LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION: TELEPHONE: (206) 224-0718
TELEPAX: (206) 224-0779
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: CHILL
ADDRESSEE: PLLC
CORRECT: 1420 Fifth Avenue, Suite 2800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc_feature
LOCATION: 1..19
OTHER INFORMATION: /product= "Primer 3.B (Table 1)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Shelton, Dennis
REGISTRATION NUMBER: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: FILING DATE:
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2662 AAAAAAAAAAAAAAAAA 2679
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                               18;
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                           0.7%; Score 18; ilarity 100.0%; Pred. No. Conservative 0; Mismatc
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100.0%; Pred. No.
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                            Mismatches
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                                            DB 3; I
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                                                         Length 19;
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APPLICANT: Cook, Phillip D
APPLICANT: Manoharan, Muthiah
APPLICANT: Kawasaki, Andrew
ITITLE OF INVENTION: Aminooxy-Modified Oligonucleotides
FILE REFERENCE: ISIS2824
CURRENT APPLICATION NUMBER: US/09/016,520A
CURRENT FILING DATE: 1998-01-30
EARLIER APPLICATION NUMBER: 60/037,143
EARLIER APPLICATION NUMBER: 60/037,143
EARLIER FILING DATE: 1997-02-14
NUMBER OF SEQ ID NOS: 47
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 20
LENGTH: 19
TYPE: NNA
                                                            US-09-016-520-21
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US-09-016-520-21/c
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US-09-016-520-20/c
, Sequence 20, Application US/09016520A
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Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/016,520A
CURRENT FILING DATE: 1998-01-30
EARLIER APPLICATION NUMBER: 60/037,143
EARLIER FILING DATE: 1997-02-14
NUMBER OF SEQ ID NOS: 47
SOFTMARE: Patentin Ver. 2.1
SEQ ID NO 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21, Application US/09016520A Patent No. 6127533
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                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Cook, Phillip D
APPLICANT: Manoharan, Muthiah
APPLICANT: Kawasaki, Andrew
TITLE OF INVENTION: Aminooxy-Modified Oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: ISIS2824
                                                                                                                                                                                                        ORGANISM: Artificial Sequence
                                                                      OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: Sequence
                                                                                                                                                 NAME/KEY: misc_feature
LOCATION: (15)..(18)
                                                                                                                   FEATURE:
                                                                                                                               OTHER INFORMATION: 5-methyl-2'-dimethylaminooxyethoxy
                                                                                                                                                                                                  FEATURE:
                                                                                                                                                                                                                                      TYPE: DNA
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LOCATION: (15)..(18)
OTHER INFORMATION: 5-methyl-2'-aminooxyethoxy
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ORGANISM: Artificial Sequence
FEATURE:
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100.0%; Pred. No.
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Score 18;
Pred. No.
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o. 6.1e+02;
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DB 3; Length 19; 6.1e+02;
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US-09-016-520-23
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US-09-016-520-23/c
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US-09-016-520-22/c
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Patent No. 6127533
GENERAL INFORMATION:
APPLICANT: Cook, Phillip D
APPLICANT: Mancharan, Muthiah
APPLICANT: Kawasaki, Andrew
TITLE OF INVENTION: Aninooxy-Modified Oligonucleotides
FILE REFERENCE: ISIS2824
                                                                                                                                                                                                              SEQ ID NO 23
LENGTH: 19
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SEQ ID NO 22
LENGTH: 19
                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/016,520A
CURRENT FILING DATE: 1998-01-30
EARLIER APPLICATION NUMBER: 60/037,143
EARLIER FILING DATE: 1997-02-14
NUMBER OF SEQ ID NOS: 47
NUMBER OF SEQ ID NOS: 47
SOFTWARE: Patentin Ver. 2.1
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APPLICANT: Cook, Ph
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  FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
OTHER INFORMATION: Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Mancharan, Muthiah
APPLICANT: Kawasaki, Andrew
TITLE OF INVENTION: Aminooxy-Modified Oligonucleotides
FILE REFERENCE: ISIS2824
                                                                                                                                             TYPE: DNA
ORGANISM: Artificial Sequence
                                                                          OTHER INFORMATION: 5-methyl-2'-dimethylaminooxyethoxy
                                                                                                     NAME/KEY: misc_feature
LOCATION: (16)..(19)
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LOCATION: (15)..(18)
OTHER_INFORMATION: 2'-methoxyethoxy
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RESULT 47
US-09-016-520-25/c
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US-09-016-520-24/c
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Best Local Sim
Matches 18;
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SEQ ID NO 25
LENGTH: 19
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GENERAL INFORMATION:
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APPLICANT: Manoharan, Muthiah
APPLICANT: Kawasaki, Andrew
TITLE OF INVENTION: Aminooxy-Modified Oligonucleotides
FILE REFERENCE: ISIS2824
CURRENT APPLICATION NUMBER: US/09/016,520A
CURRENT FILING DATE: 1998-01-30
EARLIER APPLICATION NUMBER: 60/037,143
EARLIER FILING DATE: 1997-02-14
NUMBER: OF SEQ ID NOS: 47
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CURRENT FILING DATE: 1998-01-30
EARLIER APPLICATION NUMBER: 60/037,143
EARLIER FILING DATE: 1997-02-14
NUMBER OF SEQ ID NOS: 47
SOFTWARE: Patentin Ver. 2.1
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APPLICANT: Manoharan, Muthiah
APPLICANT: Kawasaki, Andrew
TITLE OF INVENTION: Aminooxy-Modified Oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE: ISIS2824
         ORGANISM: Artificial Sequence
PEATURE:
PEATURE:
NAME/KEY: misc_feature
LOCATION: (16)...(19)
OTHER INFORMATION: 5-methyl-2'-O-propyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: Sequence
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LOCATION: (16)..(19)
OTHER_INFORMATION: 5-methyl-2'-methoxyethoxy
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ORGANISM: Artificial Sequence
                                                                                                           TYPE: DNA
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FEATURE:
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les 18; Conservative (
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100.0%; Pred. No.
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APPLICANT: Cook, Phillip D
APPLICANT: Manoharan, Muthiah
APPLICANT: Kawasaki, Andrew
ITITE OF INVENTION: Androoxy-Modified Oligonucleotides
FILE REFERENCE: ISIS2824
CURRENT APPLICATION NUMBER: US/09/016,520A
CURRENT FILING DATE: 1998-01-30
EARLIER APPLICATION NUMBER: 60/037,143
EARLIER FILING DATE: 1997-02-14
NUMBER OF SEQ ID NOS: 47
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 26
LENGTH: 19
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
NAME/KEY: misc feature
LOCATION: (18)
OTHER INFORMATION: 5-methyl-2'-dimethylaminooxyethoxy
FEATURE:
INFORMATION: 1897-02-14
CURRENT APPLICATION NUMBER: US/09/016,520A
CURRENT FILING DATE: 1998-01-30
EARLIER APPLICATION UMBER: 60/037,143
EARLIER FILING DATE: 1997-02-14
NUMBER OF SEQ ID NOS: 47
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 27
LENGTH: 19
TABE: DNA
ORGANISM: Artificial Sequence
FEATURE:
NAME/KEY: misc_feature
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US-09-016-520-26/c
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APPLICANT: Manoharan, Muthiah
APPLICANT: Kawasaki, Andrew
TITLE OF INVENTION: Aminooxy-Modified Oligonucleotides
FILE REFERENCE: ISIS2824
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OTHER INFORMATION:
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Pred. No.
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hes 0;
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US-09-016-520-33/c
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; LOCATION: (15)..(18)
; OTHER INFORMATION: 5-methyl-2'-dimethylaminooxyethoxy
US-09-016-520-31
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SOFTWARE: Pa
SEQ ID NO 33
LENGTH: 19
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Best Local Similarity 100.
Matches 18; Conservative
                                                                                                                                                                                                                                                       Sequence 33, Application US/09016520A Patent No. 6127533
                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                               APPLICANT: Cook, Phillip D
APPLICANT: Manoharan, Muthiah
APPLICANT: Manoharan, Muthiah
APPLICANT: Kawabaki, Andrew
TITLE OF INVENTION: Aminooxy-Modified Oligonucleotides
FILE REFERENCE: ISIS2824
CURRENT FILING DATE: 1998-01-30
EARLIER APPLICATION NUMBER: US/09/016,520A
CURRENT FILING DATE: 1997-02-14
NUMBER: ESCAL DATE: 1997-02-14
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GENERAL INFORMATION:
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                                                 NUMBER OF SEQ ID NOS: 47
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LENGTH: 19
TYPE: DNA
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EARLIER FILING DATE: 1997-02-14
NUMBER OF SEQ ID NOS: 47
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APPLICANT: Manoharan, Muthiah
APPLICANT: Kawasaki, Andrew
TITLE OF INVENTION: Aminooxy-Modified Oligonucleotides
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100.0%; Pred. No. 6.1
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0; Mismatches
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RESULT 53
US-09-016-520-44/c
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; LOCATION: (16)...(19)
; OTHER INFORMATION: 5-methyl-2'-dimethylaminooxyethoxy
US-09-016-520-34
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US-09-016-520-34/c
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                                                                                                                                    Sequence 44, Application US/09016520A
PATENT NO. 6127533
GENERAL INFORMATION:
APPLICANT: Cook, Phillip D
APPLICANT: Manoharan, Muthiah
APPLICANT: Kawasaki, Andrew
TITLE OF INVENTION: Aminooxy-Modified Oligonucleotides
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Best Local Similarity 100.0%; P
Matches 18; Conservative 0;
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SEQ ID NO 34
LENGTH: 19
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FILE REFERENCE: ISIS2824

CURRENT APPLICATION NUMBER: US/09/016,520A

CURRENT FILING DATE: 1998-01-30

EARLIER APPLICATION NUMBER: 60/037,143

EARLIER FILING DATE: 1997-02-14

NUMBER OF SEQ ID NOS: 47

SOPTWARE: Patentin Ver. 2.1
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Best Local Similarity
Matches 18; Conserv
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APPLICANT: Manoharan, Muthiah
APPLICANT: Kawasaki, Andrew
TITLE OF INVENTION: Aminooxy-Modified Oligonucleotides
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OTHER INFORMATION: 5-methyl-2'-dimethylaminooxyethoxy
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00.0%; Pred. No.
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APPLICANT: Kawasaki, Andrew M
APPLICANT: Kawasaki, Andrew M
APPLICANT: Cook, Phillip Dan
APPLICANT: Cook, Phillip Dan
APPLICANT: Praser, Allister S
APPLICANT: Praser, Allister S
APPLICANT: Prakash, Thazha P
ITILE OF INVENTION: 2'-O-acctamido Modified Monomers and
FILE REFERENCE: ISISA971
CURRENT APPLICATION NUMBER: US/09/378,568
CURRENT APPLICATION NUMBER: 05/09/378,568
CURRENT FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 6
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 4
LENGTH: 19
TYDER: DNA
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; NAME/KEY: misc_feature
; LOCATION: (15) ... (18)
; OTHER INFORMATION: 2'-methyleneiminooxyethoxy
US-09-016-520-44
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US-09-130-973-20/c
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US-09-378-568-4/c
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Patent No. 6172209
GENERAL INFORMATION:
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Patent No. 6147200
GENERAL INFORMATION:
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Best Local |
                                                            TITLE OF INVENTION: Aminooxy-Modified Oligonucleotides
TITLE OF INVENTION: Making Same
FILE REFERENCE: ISIS2955
CURRENT APPLICATION NUMBER: US/09/130,973
CURRENT FILING DATE: 1998-08-07
NUMBER OF SEQ ID NOS: 58
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APPLICANT: Cook, Phillip Dan
APPLICANT: Prakash, Thazha P
APPLICANT: Kawasaki, Andrew M
SOFTWARE:
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ORGANISM: Artificial Sequence
FEATURE:
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                                    PatentIn Ver. 2.1
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APPLICANT: Manue.....
APPLICANT: Cook, Phillip Dan.
APPLICANT: Cook, Phillip Dan.
APPLICANT: Prakash, Thazha P
APPLICANT: Kawasaki, Andrew M
TITLE OF INVENTION: Aminooxy-Modified Oligonucleotides And
TITLE OF INVENTION: Making Same
FILE REFERENCE: ISIS2955
CURRENT APPLICATION NUMBER: US/09/130,973
CURRENT FILING DATE: 1998-08-07
NUMBER OF SEQ ID NOS: 58
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LENGTH: 19
TYPE: DNA
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Best Local Similarity 100.0%; P
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CURRENT FILING DATE: 1998-08-07
NUMBER OF SEQ ID NOS: 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Aminooxy-Modified Oligonucleotides And Methods TITLE OF INVENTION: Making Same FILE REFERENCE: ISIS2955
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Prakash, Thazha P
APPLICANT: Kawasaki, Andrew I
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
NAME/KEY: misc feature
LOCATION: (15)...(18)
OTHER INFORMATION: 5 methyl, 2'-dimethylaminooxyethoxy
OTHER INFORMATION: Description of Artificial Sequence:
OTHER INFORMATION: Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LOCATION: (15)...(18) OTHER INFORMATION: 5 methyl, 2'-aminooxyethoxy OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
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18; Conservative 0;
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100.0%; Pred. No.
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Pred. No.
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. 6.1e+02;
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RESULT 59
US-09-130-973-24/c
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                                               APPLICANT: Manoharan, Muthiah APPLICANT: Cook, Phillip Dan APPLICANT: Prakash, Thazha P APPLICANT: Kawasaki, Andrew M TITLE OF INVENTION: Maninoxy-Moritte Reference: ISIS2955
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Best Local S
Matches 18
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                                                                                                                                                                             GENERAL INFORMATION:
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Best Local Similarity
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LENGTH: 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Prakash, Thazha P
APPLICANT: Kawasaki, Andrew M
TITLE OF INVENTION: Making Same
TITLE OF INVENTION: Making Same
FILE REFERENCE: ISIS2955
CURRENT FILING DATE: 1998-08-07
CURRENT FILING DATE: 1998-08-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
CURRENT APPLICATION NUMBER: US/09/130,973
CURRENT FILING DATE: 1998-08-07
NUMBER OF SEQ ID NOS: 58
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SOFTWARE: Patentin Ve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: misc feature
LOCATION: (16)...(19)
COTHER INFORMATION: 2'-O-dimethylaminooxyethyl
OTHER INFORMATION: Description of Artificial Sequence:
OTHER INFORMATION: Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: misc feature
LOCATION: (15)..(18)
OTHER INFORMATION: 2'-O-methoxyethyl (MOE)
OTHER INFORMATION: Description of Artificial Sequence:
OTHER INFORMATION: Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
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                                                                  Aminooxy-Modified Oligonucleotides And Methods For Making Same
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6.1e+02;
hes 0;
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RESULT 61
US-09-130-973-26/c
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APPLICANT: Mancharan, Muthiah
APPLICANT: Cook, Phillip Dan
APPLICANT: Prakash, Thazha P
APPLICANT: Kawasaki, Andrew M
TITLE OF INVENTION: Aminoxy-Modified Oligo
TITLE OF INVENTION: Making Same
FILE REFERENCE: ISIS2955
CURRENT APPLICATION NUMBER: US/09/130,973
CURRENT FILING DATE: 1998-08-07
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US-09-130-973-25/c
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; GENERAL INFORMATION:
; APPLICANT: Manchara;
; APPLICANT: Cook, Ph
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                                                                                                                                                                      Sequence 26, Application US/09130973 Patent No. 6172209
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SEQ ID NO 25
LENGTH: 19
TYPE: DNA
ORGANISM: Artificial Sequence
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Best Local Similarity
                                                                                                                                                     GENERAL INFORMATION:
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SEQ ID NO 24
LENGTH: 19
TYPE: DNA
ORGANISM: Artificial Sequence
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NAME/KEY: misc feature
NAME/KEY: (16)..(19)
LOCATION: (16)..(19)
OTHER INFORMATION: 2'-O-propyl
OTHER INFORMATION: Description of Artificial Sequence
OTHER INFORMATION: Sequence
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LOCATION: (16)..(19)
OTHER INFORMATION: 2'-O-methoxyethyl
OTHER INFORMATION: Description of Artificial Sequence:
OTHER INFORMATION: Sequence
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Kawasaki, Andrev
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Cook, Phillip Dan
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Aminooxy-Modified Oligonucleotides
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100.0%; Pred. No.
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100.0%; Pred. No.
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. 6.1e+02
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6.1e+02;
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                                                                And Methods For
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RESULT 63
US-09-130-973-31/c
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US-09-130-973-27/c
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SEQ ID NO 27
LENGTH: 19
                             GENERAL INFORMATION:
APPLICANT: Mancharan, Muthiah
APPLICANT: Cook, Phillip Dan
APPLICANT: Prakash, Thazha P
APPLICANT: Kawasaki, Andrew M
TITLE OF INVENTION: Aminooxy-Modified Oligonucleotides
TITLE DEFERENCE: TITLE OF INVENTION: Making Same
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Patent No. 6172209
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
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SOFTWARE: PatentIn Ve
SEQ ID NO 26
LENGTH: 19
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Best Local :
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APPLICANT: Kawasaki, Andrew M
TITLE OF INVENTION: Aminooxy-Modified Oligonucleotides
TITLE OF INVENTION: Making Same
FILE REFERENCE: ISIS2955
CURRENT APPLICATION NUMBER: US/09/130,973
CURRENT FILLING DATE: 1998-08-07
NUMBER OF SEQ ID NOS: 58
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FILE REFERENCE: ISIS2955
CURRENT APPLICATION NUMBER: US/09/130,973
                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
NAME/KEY: misc feature
LOCATION: (18)
OTHER INFORMATION: 5 methyl, 2'-0-methoxyethyl
OTHER INFORMATION: Description of Artificial Sequence:
OTHER INFORMATION: Sequence
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TYPE: DNA
ORGANISM: Artificial Sequence
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NAME/KEY: misc_feature
LOCATION: (18)
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100.0%; Pred. No. 6.1
tive 0; Mismatches
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GENERAL INFORMATION:
APPLICANT: Mancharar
APPLICANT: Cook, Phi
APPLICANT: Prakash,
APPLICANT: Kawasaki
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US-09-130-973-34/c
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APPLICANT: Cook, Phillip Dan
APPLICANT: Prakash, Thazha p
APPLICANT: Kawasaki, Andrew M
TITLE OF INVENTION: Aminooxy-Modified Olige
TITLE OF INVENTION: Making Same
FILE REFERENCE: ISIS2955
CURRENT APPLICATION NUMBER: US/09/130,973
CURRENT FILING DATE: 1998-08-07
NUMBER OF SEQ ID NOS: 58
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Matches 18
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NUMBER OF SEQ ID NOS: 58

SOFTWARE: PACENTIN Ver. 2.1

SEQ ID NO.31

LENGTH: 19

TYPE: DNA

ORGANISM: Artificial Sequence
FEATURE:
NAME/KEY: misc_feature
LOCATION: (15)...(18)

OTHER INFORMATION: 2'-dimethylaminooxyethyl thymidine (T-2'-DMAQE)
OTHER INFORMATION: Description of Artificial Sequence: No. 6172209el
OTHER INFORMATION: Sequence
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                                                                                                                                             Sequence 34, Appli
Patent No. 6172209
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Patent No. 6172209
FILE REFERENCE:
             TITLE OF INVENTION: Aminooxy-Modified Oligonucleotides And Methods For TITLE OF INVENTION: Making Same
                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 19

TYPE: DNA
ORGANISM: Artificial Sequence
PRATURE:
PRATURE:
PRATURE:
NAME/KEY: misc_feature
LOCATION: (16)...(19)
OTHER INFORMATION: 2'-dimethylaminooxyethyl thymidine (T-2'-DMAOE)
OTHER INFORMATION: Description of Artificial Sequence: No. 6172209el
OTHER INFORMATION: Sequence
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                                                Manoharan, Muthiah
Cook, Phillip Dan
Prakash, Thazha P
Kawasaki, Andrew M
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                                                                                                                                                                   Application US/09130973
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; Sequence 18, Application US/09292768
; Patent No. 6194185
; GENERAL INFORMATION:
; APPLICANT: Croteau, Rodney B
; APPLICANT: Lupien, Shari L
; APPLICANT: Karp, Frank
; TITLE OF INVENTION: RECOMBINANT MATERIALS AN
; TITLE OF INVENTION: LIMONENE HYDROXYLASES
; FILE REFERENCE: wsur13463
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US-09-130-973-44/c
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CURRENT FILING DATE: 1998-08-07
NUMBER OF SEQ ID NOS: 58
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 44
                                                                                                                                                                                                                                                                                                      Query Match
Best Local :
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GENERAL INFORMATION:
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CURRENT FILING DATE: 1998-08-07
NUMBER OF SEQ ID NOS: 58
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 34
LENCTH: 19
TYPE: DNA
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TITLE OF INVENTION: Making Same
FILE REFERENCE: ISIS2955
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ORGANISM: Artificial Sequence
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NAME/KEY: misc_feature
LOCATION: (16)..(19)
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Kawasaki, Andrew M
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                               AND METHODS
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US-09-477-902-20/c
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SEQ ID NO 20
LENGTH: 19
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                                                                  Matches
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Patent No. 6194598
GENERAL INFORMATION:
APPLICANT: Cook, Phillip D
                                                                                              Query Match
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APPLICANT: Kawasaki, Andrew
TITLE OF INVENTION: Aninoxy-Modified Oligonucleotides
FILE REFERENCE: ISIS2824
CURRENT APPLICATION NUMBER: US/09/477,902
CURRENT FILING DATE: 2000-01-05
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PRIOR FILING DATE: 1998-01-30
PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1997-02-14
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CURRENT FILING DATE: 1999-04-14
EARLIER APPLICATION NUMBER: 08/881,784
EARLIER FILING DATE: 1997-06-24
                                                                                                                                                      NAME/KEY: misc feature
LOCATION: (15)...(18)
OTHER INFORMATION: 5-methyl-2'-aminooxyethoxy
OTHER INFORMATION: Description of Artificial Sequence:
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                                                                                                                                                                                                                                      TYPE: DNA ORGANISM: Artificial Sequence
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LOCATION: (1)..(19)
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                                                                                                                                            INFORMATION: Sequence
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100.0%; Pr
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3ER: 09/016,520
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6.1e+02;
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RESULT 69
US-09-477-902-21/c
; Sequence 21, Application
; Patent No. 6194598

US/09477902

GENERAL INFO

INFORMATION:

Cook, Phillip

TITLE OF INVENTION: Aminooxy-Modified Oligonucleotides

Manoharan, Muthiah

Kawasaki,

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CURRENT APPLICATION NUMBER: US/09/477,902
CURRENT FILING DATE: 2000-01-05
PRIOR APPLICATION NUMBER: 09/016,520
PRIOR FILING DATE: 1998-01-30
PRIOR PELICATION NUMBER: 60/037,143
PRIOR FILING DATE: 1997-02-14
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SOFTWARE: PatentIn Vo
SEQ ID NO 21
LENGTH: 19
                                                                                                                                                                                                                                                                                                             SOFTWARE:
SEQ ID NO 22
                                                                                      Query Match
Best Local
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Patent No. 6194598
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CURRENT APPLICATION NUMBER: US/09/477,902
CURRENT FILING DATE: 2000-01-05
PRIOR APPLICATION NUMBER: 09/016,520
PRIOR FILING DATE: 1998-01-30
PRIOR APPLICATION NUMBER: 60/037,143
PRIOR FILING DATE: 1997-02-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Manoharan, Muthiah
APPLICANT: Kawasaki, Andrew
TITLE OF INVENTION: Aminooxy-Modified Oligonucleotides
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
ORGANIAM: Artificial Sequence
PEATURE:
NAME/KEY: misc_feature
LOCATION: (15)..(18)
OTHER INFORMATION: 5-methyl-2'-dimethylaminooxyethoxy
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
                                                                                                                                                  LOCATION: (15)...(18)
OTHER INFORMATION: 2'-methoxyethoxy
OTHER INFORMATION: Beecription of Artificial Sequence: Synthetic
OTHER INFORMATION: Sequence
                                                                                                                                                                                                     FEATURE:
NAME/KEY: misc_feature
LOCATION: (15)..(18)
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                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                            TYPE: DNA
                                                                                                                                                                                                                                                                                         ENGTH: 19
                                                                                    Local Similarity
                                2662 AAAAAAAAAAAAAAAAA 2679
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                                                                    Conservative
                                                            100.0%;
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0; Mismatches
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NAME/KEY: misc feature

LOCATION: (16)...(19)

OTHER INFORMATION: 5-methyl-2'-methoxyethoxy

OTHER INFORMATION: Description of Artificial

OTHER INFORMATION: Sequence

US-09-477-902-24
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      Matches
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                     Query Match
Best Local Similarity
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PRIOR APPLICATION NUMBER: 60/0
PRIOR APPLICATION DATE: 1997-02-14
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APPLICANT: Mancharan, Muthiah
APPLICANT: Kawasaki, Andrew
TITLE OF INVENTION: Aminooxy-Modified Oligonuclectides
FILE REFERENCE: ISIS2824
CURRENT APPLICATION NUMBER: US/09/477,902
CURRENT FILING DATE: 2000-01-05
PRIOR APPLICATION NUMBER: 09/016,520
PRIOR FILING DATE: 1998-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Manoharan, Muthiah
APPLICANT: Kawasaki, Andrew
TITLE OF INVENTION: Aminooxy-Modified Oligonucleotides
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PRIOR APPLICATION NUMBER: 60/0
PRIOR FILING DATE: 1997-02-14
NUMBER OF SEQ ID NOS: 47
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NUMBER OF SEQ ID NOS:
SOFTWARE: Patentin Vo
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CURRENT APPLICATION NUMBER: US/09/477,902
CURRENT FILING DATE: 2000-01-05
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NAME/KEY: misc feature
NAME/KEY: misc feature
LOCATION: (16)..(19)
OTHER INFORMATION: 5-methyl-2'-dimethylaminooxyethoxy
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
OTHER INFORMATION: Sequence
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TYPE: DNA
ORGANISM: Artificial Sequence
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TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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RESULT 71

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RESULT 73 US-09-477-902-25/c

Sequence 25, Application US/09477902 Patent No. 6194598

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RESULT 74
US-09-477-902-26/c
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SEQ ID NO 26
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SEQ ID NO 25
LENGTH: 19
TYPE: DNA
ORGANISM: Artificial Sequence
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Best Local Similarity 100.0%;
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GENERAL INFORMATION:
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CURRENT APPLICATION NUMBER: US/09/477,902

CURRENT FILING DATE: 2000-01-05

PRIOR APPLICATION NUMBER: 09/016,520

PRIOR FILING DATE: 1998-01-30

PRIOR APPLICATION NUMBER: 60/037,143

PRIOR FILING DATE: 1997-02-14
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APPLICANT: Kawasaki, Andrew
TITLE OF INVENTION: Aminooxy-Modified Oligonucleotides
                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILE REFERENCE: ISIS2824
CURRENT APPLICATION NUMBER: US/09/477,902
CURRENT FILING DATE: 2000-01-05
PRIOR APPLICATION NUMBER: 09/016,520
PRIOR FILING DATE: 1998-01-30
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NUMBER OF SEQ ID NOS: 47
SOFTWARE: PatentIn Ver.
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APPLICANT: Manoharan, Muthiah
APPLICANT: Manoharan, Andrew
TITLE OF INVENTION: Aminooxy-Modified Oligonucleotides
OTHER INFORMATION: 5-methyl-2'-dimethylaminooxyethoxy OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
NAME/KEY: misc_feature
LOCATION: (16)..(19)
OTHER INFORMATION: 5-methyl-2'-O-propyl
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
OTHER INFORMATION: Sequence
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                                                        NAME/KEY: misc feature
                                                                                                       ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                           TYPE: DNA
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1998-01-30
TMRER: 60/037,143
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100.0%; Pred. No.
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Search completed: February Job time: 257.971 secs

2, 2005, 23:37:45

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                                                                                                                  ; NAME/KEY: misc_feature
; LOCATION: (18)<sup>-</sup>
; OTHER INFORMATION: 5-methyl-2'-methoxyethoxy
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
; OTHER INFORMATION: Sequence
US-09-477-902-27
                                                                                                                                                                                                                                   PRIOR FILING DATE: 1997-02-14
NUMBER OF SEQ ID NOS: 47
SOSTWARE: Patentin Ver. 2.1
SEQ ID NO 27
LENGTH: 19
TYPE: DNA
ORGANISM: Artificial Sequence
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Matches
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Patent No. 6194598
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CURRENT APPLICATION NUMBER: US/09/477,902
CURRENT FILING DATE: 2000-01-05
PRIOR APPLICATION NUMBER: 09/016,520
PRIOR FILING DATE: 1998-01-30
PRIOR APPLICATION NUMBER: 60/037,143
                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Kawasaki, Andrew
TITLE OF INVENTION: Aminooxy-Modified Oligonucleotides
FILE REFERENCE: ISIS2824
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                                                                            Local Similarity
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                  2662 AAAAAAAAAAAAAAAAA 2679
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Similarity 100.0%; P
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                                                           0.7%; Score 18; ilarity 100.0%; Pred. No. Conservative 0; Mismatc
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Pred. No.
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6.1e+02;
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1 14 4.4 24 8 AZ411813 2 12 3.8 14 5 BQ593541 3 12 3.8 19 8 AZ650592 5 12 3.8 20 6 CF317946 6 12 3.8 20 6 CF317946 7 12 3.8 24 6 CF291636 9 12 3.8 24 8 AZ346916 9 12 3.8 25 8 AZ374695 10 12 3.8 25 8 AZ374695 11 12 3.8 27 6 CF339613 12 12 3.8 27 9 CL680176 13 12 3.8 27 9 CL680176 14 12 3.8 29 6 CF327746 15 12 3.8 29 6 CF327746 16 12 3.8 30 2 BE904656 17 13 3.4 20 8 AZ559239 18 11 3.4 20 8 AZ559239 19 11 3.4 21 8 AZ355162 20 11 3.4 22 8 AZ367672 21 11 3.4 24 8 AZ367672	Pred. No. is the number of results predicted score greater than or equal to the score of t and is derived by analysis of the total score SUMMARIES Result Query No. Score Match Length DB ID	EST: * EST: * EST: * 2: gb_est1: * 3: gb_est2: * 3: gb_est3: * 4: gb_est3: * 6: gb_est6: * 6: gb_est6: * 7: gb_est6: * 8: gb_gss2: * 9: gb_gss2: *	Minimum DB seq length: 0 Maximum DB seq length: 30 Post-processing: Listing first 150 summaries	umber of hits satisfying chosen paramete	Searched: 32822875 seqs, 18219865908 residence of the search sequence of the sequence of	Scoring table: OLIGO_NUC Gapop 60.0 , Gapext 60.0	Title: US-10-048-046-1_COPY_81_399 Perfect score: 319 Sequence: 1 gtgaatcccgatggagcggc	Run on: February 2, 2005, 15:07:30 ; 80 (with 1186	GenCore version 5.1 Copyright (c) 1993 - 2005 Con OM nucleic - nucleic search, using sw model
AZ411813 1M0185H03 BQ593541 S013408-0 AI042533 Oy06e03.x AZ662892 1M0542D15 CF917946 HD07-N0 AZ450598 1M0249H23 CF991636 14ROOT0 AZ345816 1M00872B16 AZ374695 1M0127108 AZ793814 2M0047713 CF339613 RCL105- CL680176 PRI0128a AZ458545 1M0262B15 CF327746 NACL102- BE904656 601498767 BH862998 SALK 0929 CF339443 RCL104- AZ595239 1M0407B18 AZ355162 1M0094A24 AZ367102 2M0033B03 L32031 HUMXP10C7B AZ387872 1M0147021	sted by chance to have a of the result being printed, score distribution. Description			rs: 46458	esidues		.tacagactggggatgtcatc 319	Search time 979.597 Seconds ithout alignments) 866.381 Million cell updates/sec	5.1.6 Compugen Ltd.
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73 10 3.1 24 74 10 3.1 24 75 10 3.1 24 76 10 3.1 24 77 10 3.1 24 78 10 3.1 25 81 10 3.1 25 82 10 3.1 25 83 10 3.1 25 84 10 3.1 25 85 10 3.1 26 86 10 3.1 26 87 10 3.1 26 88 10 3.1 26 91 10 3.1 26 91 10 3.1 26 91 10 3.1 26 91 3.1 26 91 3.1 26 91 3.1 26 91 3.1 26 92 10 3.1 26 93 3.1 26 94 3.1 3.1 26 95 3.1 26 96 3.1 3.1 26 97 3.1 27 98 3.1 27	100 3.1 23 100 3.1 20	52 10 3.1 21 55 10 3.1 21 55 10 3.1 21 55 10 3.1 21 55 10 3.1 21 55 10 3.1 22 60 10 10 3.1 22 60 10 10 10 10 10 10 10 10 10 10 10 10 10	49 10 3.1 20 50 10 3.1 20 51 10 3.1 20 51 10 3 1 20	46 10 3.1 20 47 10 3.1 20 48 10 3.1 20	43 10 3.1 19 44 10 3.1 20 45 10 3.1 20	40 10 3.1 19	37 11 3.4 30 38 11 3.4 30	33 2 11 3.4 28 33 31 3.4 29 3.4 20 3.4 20 3.4 20 3.4 20 3.4 20 3.4 20 3.4 20 3.4 20 3.4 20 3.4 20 3.	25 11 3.4 26 27 11 3.4 26 28 11 3.4 26 29 11 3.4 26 29 11 3.4 26 27 27 27 27 27 27 27 27 27 27 27 27 27
73 10 3.1 22 76 10 3.1 2 2 77 10 3.1 2 2 77 10 3.1 2 2 10 3.1 2 10 3.1 2 2 10 3.1 2 2 10 3.1 2 2 10 3.1 2 2 10 3.1 2 2 10 3.1 2 10 3.1 2 2 10 3.1 2 2 10 3.1 2 2 10 3.1 2 2 10 3.1 2 2 10 3.1 2 10	10 3.1 23 8 AZ34252 10 3.1 23 8 AZ34554 10 3.1 23 8 AZ756195 10 3.1 23 8 AZ95213 10 3.1 23 9 TA125F1 10 3.1 23 9 AG19821 10 3.1 23 9 AG20252 10 3.1 23 9 AG20252	52 10 3.1 21 1 AZ66333 54 10 3.1 21 7 D11772 55 10 3.1 21 8 AZ40540 56 10 3.1 21 8 AZ60730 57 10 3.1 21 8 AZ60730 58 10 3.1 21 8 AZ60720 59 10 3.1 21 8 AZ65807 59 10 3.1 22 8 BF97969 60 10 3.1 22 6 CF30999 61 10 3.1 22 6 AZ78861 62 10 3.1 22 8 AZ78861 63 10 3.1 22 9 TAZ14F0 64 10 3.1 23 9 TAZ14F0	49 10 3.1 20 8 50 10 3.1 20 8 51 10 3.1 20 9 51 10 3 1 20 9	46 10 3.1 20 8 AZ30761 47 10 3.1 20 8 AZ31134 48 10 3.1 20 8 AZ36645	43 10 3.1 19 8 AZ99487 44 10 3.1 20 6 CF28121 45 10 3.1 20 6 CF32535	40 10 3.1 10 9 CM 5001 41 10 3.1 19 1 AIG9683 41 10 3.1 19 8 AZ58910	36 11 3.4 30 8 AZ654405 37 11 3.4 30 8 BZ763848 37 11 3.4 30 9 BZ763848 38 11 3.4 30 9 AG200629	32 11 3.4 28 8 AZ95807 32 11 3.4 28 8 AZ95807 33 11 3.4 29 8 AZ80581 34 11 3.4 29 9 TA128C0 35 11 3.4 29 9 TA128C0	25 11 3.4 25 1 26 11 3.4 26 8 27 11 3.4 26 8 28 11 3.4 26 8 29 11 3.4 26 8 29 11 3.4 26 8

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        Dunn, D., Aoyagi, A.,
Islam, H., Longacre,
                            Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia; 1 (bases 1 to 24)
                                                                  Mus musculus
                                                                                                      AZ411813.1
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24 bg
1M0185H03F Mouse 10kb plasmid U
clone UUGC1M0185H03 F, genomic
                                                                                                                                                                                                                                                   musculus (house mouse)
                                                                                                    GI:10535826
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3., Mahmoud,
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AZ441188
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AA968729
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                                     Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus.
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UUGC1M library Mus musculus genomic
    Beacorn, T.,
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                                                                                                                          survey sequence.
      Meenen, E., Pedersen, T.,
                 Duval, B., Hamil, C.,
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W54502 md01b04.r1
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1 SALK 1015
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G. brucei
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6 1M0110F02
9 1M0295C02
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AJ666289
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2M0060A03
2M0139A01
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T. brucei
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T. brucei
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   Beta vulgaris
Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Caryophyllales; Amaranthaceae; Beta.
1 (bases 1 to 14)
1 (bases 1 to 14)
                                                                                                                                                                                                                                                     14;
                                                                                                                  BQ593541 14 bp mRNA linear EST 06-DEC-2002 S013408-024-026-P02-SP6 MPIZ-ADIS-024-developing root Beta vulgaris clone 024-026-P02 5-PRIME, mRNA sequence.
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  Herwig, R.,
                                                                         Beta vulgaris
                                                                                                           BQ593541
                                                                                                 BQ593541.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Class: plasmid ends
High quality sequence stop: 24.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0185 row: H Column: 03
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University of Utah
Rm. 308, Biomedical Polymers Research
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Fax: 801 585 7177
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                                                                                                                                                                                                                      CACGTCCTCCTGAG
                                                                                                                                                                                                        CACGTCCTCCTGAG
                                                                                                                                                                                                                                                     ilarity 100.0%;
Conservative 0;
                                                                                                                                                                                                                                                                                                           Laboratory Mouse DNA Resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNN42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XI10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /lab_host="B. Coli strain XL10-Gold, T1-resistant, /clone_1b="Mouse 10kb plasmid UUGCIM library" /note="Vector: pW042nv; Purified genomic DNA from Purisculus C57BL/6J (male) was obtained from the Jac
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mol_type="genomic DNA
/strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone="UUGC1M0185H03"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sex="Male"
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les 12; Conserv
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                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 18)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
                                                                                                                                                                                                                                                                                                                                                                                                               oy06e03.xl Soares senescent fibroblasts NbHSF Homo clone IMAGE:1665052 3' similar to TR:Q15662 Q15662 TRANSFORMATION-RELATED PROTEIN ;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        μ.
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Max-Planck-Institute for Plant Breeding Research
Carl-von-Linne Weg 10, 50829 Koeln, Germany
Fax: 00492215062851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and Radelof, U. Construction of a 'unigene' cDNA clone set by oligonucleotide fingerprinting allows access to 25 000 potential sugar beet genes plant J. 32 (5), 845-857 (2002) 22362189
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IMAGE Consortium (info@image.llnl.gov) for further information. Trace considered overall poor quality Seg primer: -40ml3 fwd. ET from Amersham
                                                              Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the
                                                                                                                                                                                                                                                                                                                                                                       AI042533.1 GI:3281727
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                                                                                                                                        Unpublished (1997)
                                                                                                                                                                                 National Cancer Institute, Cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SP6-Sall-CCACGCGTCCG-5prime-cDNA-polyA-CC-Notl-T7; Note: Sequencing granted in the context of the GABI-Beet project, local PI: Dr. Katharina Schneider, coordinator: Prof. Christian Jung; Sequence submission managed by RZPD/GABI-Primary database: http://gabi.rzpd.de"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="Wector: pCMVSPORT6; Site 1: Sall; Site 2: Not1; cDNA library from sugar beet, library provided by KWS Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact: b.schulz@kws.de; cloning sites Sall-Not1, primer sites and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               orientation:
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/lab_host="EMDH10B"
/clone_lib="MBIZ-ADIS-024-developing root"
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/db_xref="taxon:161934"
/clone="024-026-p02"
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/mol_type="mRNA"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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                                                                                                                                                                     source
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                                                                                                                                                                                                                                                          Tel: 801 585 5606

Fax: 801 585 7177

Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.

Plate: 0542 row: D column: 15
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       225 TGACACTGGAAG 236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AZ662892

19 bp DNA line
1M0542D15F Mouse 10kb plasmid UUGCIM library Mu
clone UUGCIM0542D15 F, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5 TGACACTGGAAG 16
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 (bases 1 to 19)
Dunn, D., Aoyagi, A., Barber, M.,
                                                                                                                                                                                                  High quality sequence stop: 19.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mouse whole genome scaffolding with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AZ662892.1 GI:11800038
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              308,
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/mol type="mcNA"
/mb xref="taxon:9806"
/db xref="taxon:9806"
/clone="IMAGE:1665052"
/tissue_type="senescent fibroblast"
/tissue_type="senescent fibroblasts NbHSF"
/clone_lib="Soares senescent fibroblasts NbHSF"
/clone_lib="Soares senescent fibroblasts NbHSF"
/clone_tib="Soares senescent fibroblasts NbHSF"
/clone="Topic Name of NbHSF"
/clone="IMAGE:1665052"
/clone="IMAGE:1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Biomedical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         double-stranded cDNA was size selected, ligated to Eccadapters (Pharmacia), digested with Not I and cloned it the Not I and Eco RI sites of a modified pT7T3 vector
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (Pharmacia). Library went through one round of normalization to a Cot = 5. Library constructed by Bento Soares and M.Fatima Bonaldo."
/db_xref="taxon:10090"
/clone="UUGC1M0542D15"
                                                               /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                  organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polymers Research Bldg.,
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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UUGC1M library Mus musculus genomic
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5.5e+05;
hes 0;
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derived from rice Histone Deacetylase overexpression

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242 AGCACCAGTGGA 253
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oryza sativa (japonica cultivar-group)
oryza sativa (japonica cultivar-group)
oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyt
Spermatophyta; Magnoliophyta; Liliopsida; Poales;
Ehrharroideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Yongin, Kyeonggi, Ko
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., L
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Nahm B.H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CF317946.1 GI:33689707
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 20)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwp2 (gil4732114 |gb|AF129072.1), a copy-number inducible derivative of plasmid Rl. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       bhnahm@ggbio.com, bhn.
Location/Qualifiers
                                                                                                                                                               /Clone lib="OsHDAC1-overexpressing transgenic rice plasmid cDNA library (HD)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"/clone_lib="Mouse_10kb_plasmid_UUGCIM_library"/note="Vector: FWD42nv; Purified_genomic_DNA_from_M.musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_f
/note="Vector: pCR4-TOPO; Site 1: EcoRI; Callus was treated with ABA(20um) for 1hr. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was
                                                                                                                                                                                                                                                                                            tissue_type="callus"
/dev_stage="proliferated callus on 2N6 media for 2 weeks"
/lab_host="E.coli DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Oryza sativa (japonica cultivar-group)"
/mol_type="mRNA"
/cultivar="Nackdong"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             clone="HD--07-N06"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            db_xref="taxon:39947"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  bhnahm@bio.myongji.ac.kr
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a; Poales; Poaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 19;
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RESULT 5 CF317946/c

DEFINITION

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ORIGIN

Matches Query Match Best Local (

TITLE JOURNAL COMMENT

FEATURES

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REFERENCE AUTHORS

SOURCE

ORGANISM

ŒYWORDS VERSION ACCESSION

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JOURNAL COMMENT
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AUTHORS
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AZ450598
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KEYWORDS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Welse, R. Mouse whole genome scaffolding with paired end reads from 10kb Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Class: plasmid ends
High quality sequence stop: 23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Err
Plate: 0249 row: H column: 2
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Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AZ450598 23 bp DNA linear GSS 04-OCT-200
1M0249H23F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0249H23 F, genomic survey sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (2000)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into
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                                                                                                                                                                                                                                                                                                                                                                                           /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/Clone lib="Mouse 10kb plasmid UUGC1M library"/note="Vector: PW042TV; Purified genomic DNA from M.musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                        laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                clone="UUGC1M0249H23"
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100.0%; Pred. No. 5.9
tive 0; Mismatches
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1 (bases 1 to 24)

Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H., Shin,Y.C., Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
                                                                AZ346816 24 bp DNA linear GSS 2: 1M0082B16F Mouse 10kb plasmid UUGC1M library Mus musculus clone UUGC1M0082B16 F, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
Email: bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
Contact: Nahm B.H.
Genomics and Genetics Institute, GreenGene Biotech Inc.; Division
Genetics and Bioinformatics, MyongJi University
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Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CF291636 24 bp mRNA linear EST 14-AUG-2003
14ROOT--02-C09.g1 Rice root plasmid cDNA library (14ROOT) Oryza
sativa (japonica cultivar-group) cDNA clone 14ROOT--02-C09, mRNA
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musculus (house mouse)
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ilarity 100.0%; Pred. No. 5.5e+05;
Conservative 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                   /clone_lib="rice root plasmid cDNA library (14ROOT)"
/note="Wector: pCR4-TOPO; Site_1: EcoRI; mRNA was capped
with oligoribonucleotides and then used as templates for
RT-PCR."
                                                                                                                                                                                                                                                                                                                                                                                                                /dev_stage="14 days after germination"
/lab_host="E.coli DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                 /cultivar="Nackdong"
/db_xref="taxon:39947"
/clone="14ROOT--02-C09"
/tissue_type="root"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Oryza sativa (japonica cultivar-group)"
/mol_type="mRNA"
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b; Pred. No. 5.5e+05;
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Mus musculus (house mouse)
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151 CCTTTCCTTCCC 162
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Contact: Robert B. I
University of Utah (
University of Utah (
University of Utah (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R. Mouse whole genome scaffolding with paired end reads from 10kb Nose, whole genome scaffolding with paired end reads from 10kb Nose, whole genome scaffolding with paired end reads from 10kb Nose, whole genome scaffolding with paired end reads from 10kb Nose, whole genome scaffolding with paired end reads from 10kb Nose, whole genome scaffolding with paired end reads from 10kb Nose, whole genome scaffolding with paired end reads from 10kb Nose, whole genome scaffolding with paired end reads from 10kb Nose, whole genome scaffolding with paired end reads from 10kb Nose, who were the scaffolding with paired end reads from 10kb Nose, who were the scaffolding with paired end reads from 10kb Nose, who were the scaffolding with paired end reads from 10kb Nose, who were the scaffolding with paired end reads from 10kb Nose, who were the scaffolding with paired end reads from 10kb Nose, who were the scaffolding with paired end reads from 10kb Nose, who were the scaffolding with paired end reads from 10kb Nose, who were the scaffolding with paired end reads from 10kb Nose, who were the scaffolding with paired end reads from 10kb Nose, who were the scaffolding with paired end reads from 10kb Nose, who were the scaffolding with paired end reads from 10kb Nose, who were the scaffolding with paired end reads from 10kb Nose, who were the scaffolding with paired end reads from 10kb Nose, who were the scaffolding with paired end reads from 10kb Nose, who were the scaffolding with paired end reads from 10kb Nose, who were the scaffolding with paired end reads from 10kb Nose, who were the scaffolding with paired end reads from 10kb Nose, who were the scaffolding with paired end reads from 10kb Nose, who were the scaffolding with paired end reads from 10kb Nose, who were the scaffolding with paired end reads from 10kb No
AZ374695
AZ374695.1 GI:10488395
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                                                                                                                                            1M0127I08R Mouse 10kb plasmid clone UUGC1M0127I08 R, genomic
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Insert Length: 10000 Std Error:
Plate: 0082 rw: B column: 16
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Fax: 801 585 7177
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/strain="C57BL/6J"
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                                                                                     AZ793814 25 bp DNA linear GSS it 2M0047J13F Mouse 10kb plasmid UUGCIM library Mus musculus clone UUGC2M0047J13 F, genomic survey sequence.
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GSS.
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Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
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Seq primer: CACAAAGGAAACAGCTATGACC
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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Contact: Robert B. Weiss
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/note="Vector: PWD42nv, burified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackso
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/strain="C57BL/6J"
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Contact: Robert B.
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Fax: 801 585 7177
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Rm. 308, Biomedical Polymers Research Bldg.,
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Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
Email: ralf.sommer@tuebingen.mpg.de
This library was generated at Caltech,
sequenced at Vancouver, Canada.
                                                                                                         Max-Planck-Institute for Developmental Biology Spemannstr. 37-39, Tuebingen D-72076, Germany
                                                                                                                                                                                            Srinivasan,J., Otto,G.W., Kahlow,U., Geisler,R. and Sommer,R.J. AppaDB: an AcedB database for the nematode satellite organism Pristionchus pacificus
RULLELIC Acids Res. 32 (1), D421-D422 (2004)
                                                                                                                                                                                                                                                                                                                          Pristionchus pacificus
Eukaryota; Metazoa; Nematoda; Chromadorea; Diplogasterida;
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CL680176.1 GI:50187006
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Fax: 82 31 321 6355
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                                                                                                                                                       Evolutionary Biology
                                                                                                                                                                              Contact: Sommer RJ
                                                                                                                                                                                                                                                                                                          Neodiplogasteridae, Pristionchus.
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                                                                                     Tel: 00497071601371
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Location/Qualifiers
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/mol_type="mRNA"
/culTivar="Nackdong"
/db_xref="taxon:39947"
/clone="RCL1--05-G12"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="Vector: pBluescript SK(+); Site 1: SstI; Site 2: XhOI; cDNA was inserted into landa Uni-ZAP XR vector at 5' end with SstI and 3' end with XhOI site. Callus was induced on 2N6 media for 30 days and cultured for 36hrs on regenerated media"
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Pred. No. 5.5e+05;
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Pax: 801 585 7177

Email: ddunn@genetics.utah.edu

Insert Length: 10000 Std Error: 0.1

Plate: 0262 row: B column: 15

Seg primer: CACCAGGAAACAGCTATGACC

Class: plasmid ends
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Contact: Robert B. I
University of Utah (
University of Utah)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus (bases 1 to 28)
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clone UUGC1M0262B15 R, genomic
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(http://www.jax.org/resources/documents/dnares/). The DN7 was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant DN7 by the sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                        /lab host="E. Coli strain XL10-Gold, T1-resistant, Clone lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: pWD42Tnv; Purified genomic DNA from lmusculus C57BL/6J (male) was obtained from the Jac
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                                                                                                                   Laboratory Mouse DNA Resource
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/mol_type="genomic DNA"
/strain="California"
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                                                                                                                                                                                                                                                              db_xref="taxon:10090"
                                                                                                                                                                                                                                                                              strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                 /mol_type="genomic DNA"
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                                                                                                                                                                                                                    sex="Male"
                                                                                                                                                                                                                                           clone="UUGC1M0262B15"
                                                                                                                                                                                                                                                                                                                     organism="Mus musculus"
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Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Enrhantoideae; Oryzeae; Oryza.
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H.,
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
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NACL--02-F18.gl Rice callus plasmid cDNA library (NACL) Oryza
Sativa (japonica cultivar-group) cDNA clone NACL--02-F18, mRNA
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82 31 330 6193
82 31 321 6355
                                                                                                                                    Conservative
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Location/Qualifiers
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                                                                                                                                                                                                                                              /clone_lib="Rice callus plasmid cDNA library (NACL)"
/note="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA was capped
/nith_nigoribonucleotides and then used as templates for
                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Oryza sativa (japonica
/mol_type="mRNA"
/cultivar="Nackdong"
                                                                                                                                                                                                                                                                                                            dev_stage="proliferated callus on 2N6 media for 30 days"
/lab_host="E_coli DH10B"
                                                                                                                                                                                                                                                                                                                                                                         /db_xref="taxon:39947"
/clone="NACL--02-F18"
                                                                                                                                                                                                                                                                                                                                                     tissue_type="callus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GI:33803742
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Pred. No.
                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          bhnahm@bio.myongji.ac.kr
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                                                                                                                          DB 6; Lc.
3. 5.5e+05;
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5.5e+05;
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Best Local
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Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
                                                                                                                                  Alonso, J.M., Leisse, T.J., Barajas, P., Chen, H., Cheuk, R., Gadrinab, C., Jeske, A., Karnes, M., Kim, C.J., Parker, H., Prednis, Shinn, P., Zimmerman, J. and Ecker, J.R.
                                                                                                                                                                                                                                                                                                                                                                                        BH862998 30 bp DNA linear GSS 05-AUG-200
SALK_092934 Arabidopsis thaliana TDNA insertion lines Arabidopsis
thaliana genomic clone SALK_092934, genomic survey sequence.
BH862998
                                                                        Arabidopsis Genome 
Unpublished (2001)
                                                                                                                                                                                                                            Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                Arabidopsis thaliana (thale cress)
Arabidopsis thaliana
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
                                                                                                                 A Sequence-Indexed Library of Insertion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             http://image.llnl.gov
Plate: LLAM9700 row: e column: 18
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mRNA sequence.
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601498767F1 NIH_MGC_70 Homo sapiens cDNA clone IMAGE:3900665 5',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note="Organ: pancreas; Vector: pCMV-SPORT6; Site 1: NotI; Site 2: SalI; Cloned unidirectionally. Primer: Oligo dT. Average insert size 1.1 kb. Library constructed by Life Technologies."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="teaxon:9606"
/clone="IMAGE:3900665"
/clone="EDH10B (phage-resistant)"
/lab host="DH10B (phage-resistant)"
/clone=_lb="NIH_MGC_70"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
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end with XhoI site. Callus was for 30 days and cultured for 36hrs on

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Best Local
Matches
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                                                                                                                                                                                                                                                                                                                     Yongin, Kyeonggi, Ko
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                          Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeonggi, Korea
                                                                                                                                                                                                                                                                                                                                                                                                                                               1 (bases 1 to 20)

Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H., Shin,Y.C.,

Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnollophyta, Liliopsida, Poales, Poaceae,
Spermattoideae, Oryzeae, Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CF339443 20 bp mRNA 1
RCL1--04-003.gl Regenerated callus lambda ph
Oryza sativa (japonica cultivar-group) cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Nahm B.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CF339443.1 GI:33827271
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This is single pass sequence recovered from the left TDNA. This sequence lies within 300 bases of the 5' \epsilon
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TDNA tagged
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone lib="Arabidopsis thaliana TDNA insertion lines"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
                                                                                        /tissue_type="callus"
/dev_stage="proliferated callus on
/lab_host="E.coli SOLR"
                                                                                                                                                                                          organism="Oryza sativa
/mol_type="mRNA"
/cultivar="Nackdong"
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/ecotype="Col-0"
        note="Vector:
(hoI; cDNA was
                                                                                                                                                                        db_xref="taxon:39947"
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                                                          elib="Regenerated callus lambda phage cDNA library
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pBluescript SK(+); Site_1: SstI; Site_2: inserted into lamda Uni-ZAP XR vector at
                                                                                                                                                                                                                                                                                                  bhnahm@bio.myongji.ac
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 8; Le
                                                                                                                                                                                                                                     (japonica cultivar-group)"
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                                                                                                              2N6 media for 30 days"
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end of
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L1--04-003,
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ACCESSION VERSION KEYWORDS

ORGANISM

FEATURES

sheared, adaptored mouse DNA was

annealed to

COMMENT

TITLE

JOURNAL

REFERENCE

AUTHORS

RESULT 17 CF339443/c LOCUS

DEFINITION

믕 S ORIGIN

FEATURES

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Unpublished (2000)
Contact: Robert B. V
University of Utah (
University of Utah
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro.
Plate: 0407 row: B column: 18
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Mouse whole genome scaffolding with paired end reads from 10kb
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2 (bases 1 to 20)
2 (bases 1 to 20)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Seq primer: CACACAGGAAACAGCTATGACC Class: plasmid ends
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus
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was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gi|4732114|gb|R7129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          end with SstI and 3' induced on 2N6 media regenerated media"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Biomedical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /lab host="E. Coli strain XL10-Gold, T1-resistant, F. /clone_lib="Mouse 10kb plasmid UUGC1M library" /note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone="UUGC1M0407B18"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sex="Male"
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100.0%; Pr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 1
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1.9e+06;
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KEYWORDS

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SOURCE
ORGANISM
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AZ355162
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AUTHORS
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        source
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von
Niederhausern,A. and Wright,D., Weiss,R.
Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19 GGCTCCTGCGT 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std 1274
Plate: 0094 row: A column: 24
Seq primer: CACACAGGAAACAGCTATGACC
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GSS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        High quality sequence stop: 21.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AZ355162 21 bp DNA linear GSS 02-OCT-20
1M0094A24R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0094A24 R, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               plasmid inserts
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           (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pDM42 (gi |4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     USA
                                                                                                                                                                                                                                                                                                            /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone_lib="Mouse_10kb plasmid_UUGCIM_library"
/note="Vector: PWD42nv; Purified_genomic_DNA_from_M.
musculus_C57BL/6J_(male)_was_obtained_from_the_Jackso
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                 Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                             clone="UUGC1M0094A24"
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1.9e+06;
thes 0;
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                                                   RESULT 21
AZ360666
LOCUS AZ360666 22 bp DNA linear GSS 02-OCT-200
DEFINITION 1M0104F04F Mouse 10kb plasmid UUGCIM library Mus musculus genomic
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COMMENT

TITLE

FEATURES

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y Match
3.4%; Score 11; DB 9; L
Local Similarity 100.0%; Pred. No. 1.9e+06;
                                                                               301 АСАСАСТССС 311
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Park, H., Kim, Y., Kim, S., Han, Y., Woo, T., Park, K., Eun, C.J., Hoon, S.T., Chu, M., Kim, H., Joo, S., Kim, C., Song, W. and Yoo, H. Direct Submission
Submitted (07-JAN-2002) Hong-Seog Park, Korea Research Institute of Bioscience and Biotechnology (KRIBB), Genome Research Center (GRC); 52, Oun-dong, Yusong-gu, Daejeon 305-333, Korea (E-mail:redstone@mail:kribb.re.kr, URL:http://phs.grc.kribb.re.kr/, Tel. 100 Apr. 100 Ap
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         of clone tracking errors. PRIMERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Park,H., Kim,Y., Kim,S., Han,Y., Woo,T., Park,K., Eun,C.J., Hoon,S.T., Chu,M., Kim,H., Joo,S., Kim,C., Song,W. and Yoo,H. BAC end sequences of Library RP-43
Unpublished
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Pan troglodytes
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Pan.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AG202462.1 GI:45234637
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                                                                                                                                                              Conservative
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R.Site 1 : EcoRI
R.Site 2 : EcoRI.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /mol_type="genomic DNA"
/db_xref="taxon:9598"
/clone="RP43-085014.T7"
                                                                                                                                                                                                                                                                                                                                                                    (cell_type="lymphocytes"
/clone_lib="RP-43 Chimpanzee Male BAC Library"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          organism="Pan troglodytes"
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GSS 02-OCT-2000

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RESULT 22
AZ787102/c
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AZ787102 22 bp DNA linear GSS 16-FEB-200 2M0033B03F Mouse 10kb plasmid UUGCIM library Mus musculus genomic
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Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0104 row: F column: 04
Seg primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         University of Utah Genome Center University of Utah Rm. 308, Biomedical Polymers Res
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished (2000)
Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weise,R. Mouse whole genome scaffolding with paired end reads from 10kb Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 22)
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Location/Qualifiers
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                                                                                                                                                                AGTGGATGAAA 13
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                                                                                                                                                                                                                                                      3.4%; Score 11; DB 8; L
(larity 100.0%; Pred. No. 1.9e+06;
Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                  (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pW042 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. col1 XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Biomedical Polymers Research Bldg., 20
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/strain="C57BL/6J"
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                                                                                RESULT 23
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       DEFINITION HUMXP10C7B Human placenta Homo sapiens cDNA clone XP2E8B, mRNA
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Rm. 308, Bio
To USA
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mRNA

EST 02-AUG-1995

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187 TCACTGTAGAA 197
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0033 row: B column: 03
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University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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AZ787102.1 GI:12925528
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Fax: 801 585 7177
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1 (bases 1 to 22)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         quality sequence stop: 22
                                                            Conservative
                                                                                                                                                                                                    was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNAsmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone_lib="Mouse_10kb_plasmid_UUGCIM_library"
/note="Vector: pWD42nv; Purified_genomic_DNA from M.
musculus_C57BL/6J (male) was obtained from the Jackson
Laboratory_Mouse_DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="Mus musculus'
/mol_type="genomic DNA"
/strain="C57BL/6J"
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                                                                                        3.4%;
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Rodentia;
                                                                                        Score 11; DB 8; L; Pred. No. 1.9e+06;
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L32031
L32031.1 GI:927073
EST.
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                                                   Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0147 row: O column: 21
Seg primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                 University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 24)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AZ387872 24 bp DNA linear GSS 02-OCT-2001 100147021R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0147021 R, genomic survey sequence.
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                                                                                                                                                                                                                                    Contact: Robert B.
                                                                                                                                                                                                                                                    Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus
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Coolbaugh,M.I., Chinault,C.A., Baldini,A., Lindsay,E.A., Zhao,Z.-Y.
and Caskey,C.T.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 24)
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96090257
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lass: plasmid ends
igh quality sequence stop: 24.
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801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /clone_lib="Human placenta"
/note="Arrayed cDNAs and cosmid libraries from human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/map="%q12-q13.1"
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           Trace considered overall poor quality Insert Length: 704 Std Error: 0.00 Seq primer: -40ml3 fwd. ET from Amersham High quality sequence stop: 1.
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AA911748 25 bp mRNA linear EST 24-AUG-1 oi15g11.s1 NCI CGAP GC4 Homo sapiens cDNA clone IMAGE:1476644 3' similar to TR:Q99628 Q99628 SIAH BINDING PROTEIN 1 ;, mRNA
                                                                                                                                                                                                                         Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher A. Moskaluk, M.D.,
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 25)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

NAtional Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                   cDNA Library Preparation: M. Bento Soares, Ph.D.
CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pw12 (gi|473114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (http://www.jax.org/resources/documents/dnares/).
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/mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
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100.0%; Pred. No. 1.9e+06;
ive 0; Mismatches 0;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                               Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0016 row: N column: 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 plasmid inserts
Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. Mouse whole genome scaffolding with paired end reads from 10kb Mouse whole genome scaffolding with paired end reads
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 26)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AZ309743 26 bp DNA linear GSS 29-SEP-20 1M0016N05F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0016N05 F, genomic survey sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Tel: 801 585 5606
Fax: 801 585 7177
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AZ309743.1 GI:10351041
                                                                                                                                                                                                                                                                                                                                           Seq primer: CGTTGTAAAACGACGGCCAGT
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                                                                                                                                                                                                                                                              s: plasmid ends
quality sequence stop: 26.
Location/Qualifiers
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//Clone lib="NCI CGAP GC4"
//note="Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; lst strand cDNA was prepared from 3 pooled germ cell tumors, and was then primed with a Not I oligo(dT) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library is normalized. Library was constructed by Bento Soares and M. Fatima Bonaldo. "
  /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"/Clone_lib="Mouse 10kb plasmid UUGCNM library"/notee="Vector: PW042nv; Purified genomic DNA from M.musculus C57BL/6J (male) was obtained from the Jacksor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Biomedical Polymers Research Bldg., 20 S. 2030
                                                                                                                     /mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0016N05"
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/tissue_type="pooled germ cell tumors"
/lab_host="DH10B"
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/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                      organism="Mus musculus"
                                                                                                 sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3.4%; Score 11; DB 1; L
100.0%; Pred. No. 1.9e+06;
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                                                                                                                                                                                                                                                                                                                                                                                     Std Error: 0.00
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from the Jackson
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26 bp DNA linear GSS 16-FEB-20
2M0045N21R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC2M0045N21 R, genomic survey semience
AZ792942
                                                                                                                                                                                                                                                                                                                                                      Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0045 row: N column: 21
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Ha
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,
Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von
Niederhausern,A. and Wright,D.,Weiss,R
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert B. Weiss University of Utah Genome University of Utah
                                                                                                                                                                                                                                                                                                      High quality sequence stop: 26.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tel: 801 585 5606
Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mouse whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
                     /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"/clone_lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: PW9270v; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Biomedical Polymers Research Bldg.,
Laboratory Mouse DNA Resource
                                                                                                                                                                                                organism="Mus musculus"
/mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                /db_xref="taxon:10090"
/clone="UUGC2M0045N21"
                                                                                                                           sex="Male"
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hes 0;
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                          Jackson
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGNAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Arabidopsis Genome Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R., Gadrinab,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., P. Shinn,P., Zimmerman,J. and Ecker,J.R.

A Sequence-Indexed Library of Insertion Mutations in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BH863411 26 bp DNA linear GSS 05-AUG-200
SALK 093836 Arabidopsis thaliana TDNA insertion lines Arabidopsis
thaliana genomic clone SALK_093836, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Class: TDNA tagged.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This is single pass sequence recovered from the left border of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Arabidopsis thaliana
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ecker@salk.edu
/clone="SALK_093836"
/clone lib="Arabidopsis thaliana TDNA insertion lines"
/clone lib="Arabidopsis thaliana lines"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
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                                                                                                                                                                                                                                                                                                                    ecotype="Col-0"
                                                                                                                                                                                                                                                                                                                                                                                  organism="Arabidopsis thaliana"
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                                                                                                                                                                                                                                                                                        xref="taxon:3702"
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                                                                                                                                                          TA45F03P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
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A Sequence-Indexed Library of Insertion Mutations in the
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/ecotype="Col-0"
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/clone="SALK_134931.24.10.x"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AI78671.0 28 bp mRNA linear EST 02-JUI uj56b01.x1 Sugano mouse liver mlia Mus musculus cDNA clone IMAGE:1923913 3' similar to SW:FHIM RAT P13437 3-KETOACYL-COA THIOLASE MITOCHONDRIAL ;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.E., Rajandream,M.A. and Barrell,B.G. Direct Submission Submitted (10-DEC-2000) Trypanosoma brucei genome Sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton,
           Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                          Contact: Marra M/WashU-NCI Mouse EST Project 1999 Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. Louis, MO
                                                                                                                                                                                            Marra,M., Hillier,L., Kucaba,T., Martin,J., Beck,C., Wylie,T., Underwood,K., Steptce,M., Theising,B., Allen,M., Bowers,Y., Person,B., Swaller,T., Gibbons,M., Pape,D., Harvey,N., Schurk,FRitter,E., Kohn,S., Shin,T., Jackson,Y., Cardenas,M., McCann,R. Waterston,R. and Wilson,R.
The WashU-NCI Mouse EST Project 1999
Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger Centre are available
at http://www.sanger.ac.uk/Projects/T_brucei/.
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|mol_type="genomic DNA"
|strain="TREU927"
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100.0%; Pred. No. 1.9e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                      AUTHORS
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                                                                                                                                                                                                              Unpublished (2000)
Contact: Robert B. W
University of Utah G
University of Utah
Rm. 308, Biomedical 1
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                                  Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 025 row: 0 column: 10
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
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                                                                                                                                                 801 585 5606
801 585 7177
quality sequence stop: 28 Location/Qualifiers
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/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone="IMAGE:1923913"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     organism="Mus musculus"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          db_xref="taxon:10090"
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100.0%;
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                                                                                                                                                                                                                                                              Genome Center
                                                                                                                                                                                                                                                                                   Weiss
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Trace considered overall poor quality
Possible reversed clone: similarity on wrong strand
                                                                                                                                                                                /clone lib="Sugano mouse liver mlia"
/note="Organ: liver; Vector: pMEL8S-F13; Site 1: DraIII (CACCATGTG); Site 2: DraIII (CACCATGTG); 1st Strand cDNA was primed with an oligo(dT) primer [ATGTGGCCTTTTTTTTTTTTTTTTTTTT]; double-stranded cDNA was ligated to a DraIII adaptor [TGTTGGCCTACTGG], digested and cloned into distinct DraIII sites of the pMEL8S-FL3 vector (5' site CACCTGTGG, 3' site CACCATGTG). XhoI shou be used to isolate the cDNA insert. Size selection was performed to exclude fragments <1.5kb. Library
constructed by Dr. Sumio Sugano (University of Tokyo Institute of Medical Science). Custom primers for sequencing: 5' end primer CTTCTGCTCTAAAAGCTGCG and 3' end primer CGACCTGCAGCTCGAGCACA."
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DB 1; Length 28; 0 Gaps 0

AZ958079 28 bp DNA line 2M0225010F Mouse 10kb plasmid UUGC2M library Mu Clone UUGC2M0225010 F, genomic survey sequence. Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus op DNA linear GSS 27-APR-200 UUGC2M library Mus musculus genomic GSS 27-APR-2001 Murinae; Mus

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostor Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Nobases T to 28)

1 (Dases T to 28)

Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R.

Nouse whole genome scaffolding with paired end reads from 10kb Mouse whole genome scaffolding with paired end reads Polymers Research Bldg., 0.00 20 <u>د</u> 2030 ; ; SLC,

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REFERENCE
AUTHORS
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VERSION
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AZ805819
              FEATURES
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                                                        Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0067 row: K column: 19
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                   Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                 Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. Mouse whole genome scaffolding with paired end reads from 10kb Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AZ805819 29 bp DNA linear GSS 20-FBB-200
2M0067K19F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC2M0067K19 F, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                   plasmid inserts
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AZ805819.1 GI:12966630
                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (2000)
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                                 quality sequence stop: 29
                                                                                                                                                                                            801 585 5606
801 585 7177
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                                                                                                                                                                                                                                                          USA
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/Clone lib="Mouse 10kb plasmid UUGCZM library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6U (female) was obtained from the Jackson
Laboratory Mouse DNA Resource
        Location/Qualifiers
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/mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3.4%; Score 11; DB 8; 100.0%; Pred. No. 1.9e+0 tive 0; Mismatches
                                                                                                                                       Std Error: 0.00
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thes 0;
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                                                                                                             to give a tight size distribution (
4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B.
                                                                                                                                                                                                                                                                                                                                       Direct Submission
Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing
project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton,
Cambridge CB10 18A, B-mail: barrell@sanger.ac.uk and
nhl@sanger.ac.uk
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Trypanosoma.

1 (bases 1 to 29)

1 (bases 1 to 29)

Hall, N., Bowman, S., Lennard, N.J., Doggett, J., Atkin, R.,
Chillingworth, C., Ormond, D., Harris, B., El-Sayed, N., Hou, L.,
Melville, S.E., Rajandream, M.A. and Barrell, B.G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     79 CGCGGAGGAGG 89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11;
           Details of T. brucei sequencing at the Sanger Centre are available at http://www.sanger.ac.uk/Projects/T_brucei/.
                                                                                                                                                                                                                                                             Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned populat Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanicall
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                                                                         Email: nelsayed@tigr.org
                                                                                                  Barrell, Oxford University Press, 1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Trypanosoma brucei
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musculus C57BL/6J (male) was obtained from the Jackson
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/strain="C57BL/6J"
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"Vector: PWD42nv; Purified genomic DNA from M.
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A clone
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                                                                                                                                                                                                                                                                                         a cloned population
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RESULT 36
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                                                                                                                                                             24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned population of Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically sheared to give a tight size distribution (4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotyenter, J.C. (Making small insert libraries for whole genome shotyenter). Waldin and B. Barrell, Oxford University Press, 1999).
AZ654405 30 bp DNA linear GSS 14
1M0528KO6R Mouse 10kb plasmid UUGC1M library Mus musculus
clone UUGC1M0528K06 R, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger Centre
at http://www.sanger.ac.uk/Projects/T_brucei/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Direct Submission
Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing
project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton,
Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 29)
Hall, N., Bowman, S., Lennard, N.J., Doggett, J., Atkin, R.,
Chillingworth, C., Ormond, D., Harris, B., El-Sayed, N., Hou, L.,
Melville, S.E., Rajandream, M.A. and Barrell, B.G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GSS.
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Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
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AL479008.1 GI:11844887
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                                                                                                                                                                                                    TCCCCAGCAAT 169
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/mol type="genomic DNA"
/strain="TREU927"
/db_xref="taxon:5691"
/clone="128c02"
                                                                                                                                                                                                                                                                                                                                                      /organism="Trypanosoma brucei"
/mol type="genomic DNA"
/strain="TREU927"
/db xref="taxon:5691"
/clone="217f07"
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tive 0; Mismatches
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100.0%; Pred. No. 1.9e+06;
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                                                                                          RESULT 37
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                                                                                                                                                         18 CCAGCACCAGT 8
                                                                                                                                                                                                                                              11;
BZ763848 30 bp DNA linear GSS 13-MAR-2003 SALK 122613.25.20.x Arabidopsis thaliana TDNA insertion lines Arabidopsis thaliana genomic clone SALK_122613.25.20.x, genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0528 row: K column: 06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (2000)
Contact: Robert B. Weiss
University of Utah
University of Utah
Rm. 308, Biomedical Polyme
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                High quality sequence stop: 30.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AZ654405.1 GI:11791551
GSS.
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Mammalia; Eutheria;
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801 585 7177
                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                             was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /lab_host="E. Coli strain XII0-Gold, T1-resistant, F-" /clone lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: pW042nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (http://www.jax.org/resources/documents/dnares/). The DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /db_xref="taxon:10090"
/clone="UUGC1M0528K06"
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/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       organism="Mus musculus"
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00.0%; Pred. No. 1.9e+06;
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Rodentia;
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AG200629
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                                                                                                                                                                                                                                                                                                     sequence.
AG200629
AG200629.1
Submitted (07-JAN-2002) Hong-Seog Park, Korea Research Institute of
                        2 (bases 1 to 30)
Park, H., Kim, Y., Kim, S., Han, Y., Woo, T., Park, K., Eun, C.J.,
Hoon, S.T., Chu, M., Kim, H., Joo, S., Kim, C., Song, W. and Yoo, H.
Direct Submission
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Park, H., Kim, Y., Kim, S., Han, Y., Wo
                                                                                                                                                                                                                                                                                      GSS.
                                                                                                                                                                                                                                                                                                                                                        Pan troglodytes DNA, clone: RP43-082L13.TJ, genomic survey
                                                                                                                          BAC end sequences of Library RP-43
                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Pan.
                                                                                                                                                                                                                                             Pan troglodytes (chimpanzee)
Pan troglodytes
                                                                                                                                                                                                                                                                                                                                                                                     AG200629
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Arabidopsis Genome 
Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R., Gadrinab,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L., Shinn,P., Zimmerman,J. and Ecker,J.R.

A Sequence-Indexed Library of Insertion Mutations in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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BZ763848
BZ763848.1 GI:28936401
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Arabidopsis thaliana
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Joseph R. Ecker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3.4%; Score 11; DB 8; Le larity 100.0%; Pred. No. 1.9e+06; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ecker@salk.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /clone="SALK 122613.25.20.x"
/clone=lib="Ārabidopsis thaliana TDNA insertion lines"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /ecotype="Col-0"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /mol_type="genomic DNA"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ocation/Qualifiers
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                                                                                                                                         Woo, T., Park, K., Eun, C.J., Kim, C., Song, W. and Yoo, H.
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CL438015
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151 CCTTTCCTTCC 161
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Class: Gene Trap.
Location/Qualifiers
                                                                                                                                                                                                                                                                                           Email: hickegg@cc.umanitoba.ca
U3NeoSVI gene trap. Tag generated by plasmid rescue. Additional
sequence information and target gene cloning can be generated. ES
cell line harboring insertion mutation of target gene is available.
Sequence analysis available from
http://140.193.242.7/esdb/public_search_frame.php?PST=PST6661-NR.Se
                                                                                                                                                                                                                                                                                                                                                                                                                                                           www.BScells.ca
Unpublished (2002)
Contact: Hicks GG
Contact: Hicks GG
Manmalian Functional Genomics Centre
Manitoba Institute of Cell Biology, Universitry of Manitoba
ON5029, 675 McDermot Ave, Winnipeg, MB R3E 0V9, Canada
Tel: 204 787 2133
Fax: 204 787 2190
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PST6661-NR.Seq MICB1 Mus musculus genomic clone PST6661-NR.Seq similar to Eif4ebp1, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hicks, G.G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
1 (bases 1 to 10)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CL438015
CL438015.1 GI:45574155
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mus musculus (house mouse)
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R.Site 1 : EcoRI
R.Site 2 : EcoRI.
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cell_type="Embryonic stem cell"
cell_line="D3H (J1 subclone)"
                                                     sex="Male"
                                                                                                                            mol_type="genomic DNA"
strain="129 sv"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /mol_type="genomic DNA"
/db_xref="taxon:9598"
/clone="RP43-082L13.TJ"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Pan troglodytes"
/mol type="genomic DNA"
                                                                            clone="PST6661-NR.Seq"
                                                                                                                                                                               organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /cell_type="1ymphocytes"
/clone_lib="RP-43 Chimpanzee Male BAC Library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /sex="male"
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                                                                                                       db_xref="taxon:10090"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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RESULT 41
AZ589109/c
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                                                                                                                                                                                    81 CGGAGGAGGG 90
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10,
19 bp DNA linear GSS 13 1M0397D22R Mouse 10kb plasmid UUGCIM library Mus musculus clone UUGC1M0397D22 R, genomic survey sequence. AZ589109
                                                                                                                                                    N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: cgapbs-r@mail.nih.gov
Life Technologies catalog #: 11548-013
Life Technologies catalog #: 11548-013
DNA Sequencing by: Washington University Genome
Clone distribution: NCI-CGAP clone distribution
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 19)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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Unpublished (1997)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert Strausberg, Ph.D.
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primer: -40UP from Gibco
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                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Homo sapiens"
/mol type="mRNA"
/db_xref="tracen:9606"
/clone="IMAGE:2324392"
/tissue type="adenocarcinoma"
/lab_host="DH108"
                                                                                                                                                                                                                                                                                                           /clone_lib="NCI_CGAP_Pan1"
/note="Organ: pancreas; Vector: pCMV-SPORT6; Site 1: Sal1;
Site_2: NotI; Cloned unidirectionally. Primer: oligo dT.
Average insert size 1.72 kb. Life Technologies catalog #:
11548-013"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clone_lib="MICB1"
/note="Vector: U3NeoSV1"
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100.0%; Pred. No. 6.
ative 0; Mismatches
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ches 0;
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Rm. 308, Bic
Rm. 308, Bic
Rm. 308, Bic
                 AZ989459 19 bp DNA linear 2M0272M17R Mouse 10kb plasmid UUGC2M library Mus clone UUGC2M0272M17 R, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0397 row: D column: 22
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R
AZ989459
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Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                      (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pW942 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/mol_type="genomic DNA"
/strain="C57BL/6J"
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AZ994873

19 bp DNA linear GSS 27-APR-2002M0280D02R Mouse 10kb plasmid UUGC2M library Mus musculus genomic clone UUGC2M0280D02 R, genomic survey sequence.
AZ994873
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Brror: 0.00
Plate: 0272 row: M column: 17
Seq primer: CACACAGGAAACAGCTATGACC
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Fax: 801 585 7177
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plasmid inserts
Unpublished (2000)
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Location/Qualifiers
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                                                                                                                                                            Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynuclectide kinase. Adaptor oligonuclectides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNDA2 (gi|4732114|gb|AFT290721), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/clone_lib="Mouse_10kb_plasmid_UUGC2M_library"
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/strain="C57BL/6J"
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20 bp mRNA linear EST 14-AUG-20 14ETL--08-C23.gl Rice etiolated leaf plasmid cDNA library (14ETL) Oryza sativa (japonica cultivar-group) cDNA clone 14ETL--08-C23, mRNA sequence.
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High quality sequence stop: 19.
Location/Qualifiers
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Insert Length: 10000 Std Error: 0.00
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20
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Contact: Robert B.
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Fax: 801 585 7177
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/clone_lib="Mouse 10kb plasmid UUGC2M library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
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/strain="C57BL/6J"
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ol_type="qenomic DNA"
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                                                                                                                                                                                                                                                                                                                              Score 10;
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0. 6.5e+06;
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Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                    JMT1--03-A01.g1 AtJMT-overexpressing transgenic rice lambda phay cDNA library (JMT1) Oryza sativa (japonica cultivar-group) cDNA clone JMT1--03-A01, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10;
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Contact: Nahm B.H.

Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeonggi, Korea
                                                                                                                                                                   1 (bases 1 to 20)

Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., Le Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H. Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                 CF325351
CF325351.1 GI:33798984
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
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1 (bases 1 to 20)

Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H.,
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
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Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
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CF281215.1
                                                                                                     Genomics and Genetics Institute, of Bioscience and Bioinformatics,
                                           Yongin, Kyeonggi, Ko
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                Contact: Nahm B.H
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bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr.
Location/Qualifiers
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA was capped
with oligoribonuclectides and then used as templates for
RT-PCR."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /dev_stage="14 days after germination"
/lab_host="8.coli DH10B"
/clone_lib="Rice etiolated leaf plasmid
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/clone="14ETL--08-C23"
/tissue_type="leaf"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism="Oryza sativa (japonica cultivar-group)"
/mol_type="mRNA"
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100.0%; Pred. No. 6.5e+06
ive 0; Mismatches 0
                                                                                      Korea
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                                                                                                       GreenGene Biotech Inc.; , MyongJi University
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                                                                                                                                                                                                                                                                                                           Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0009 row: G column: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert B. Weiss
University of Utah Genome
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B. Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Peder Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., Niederhausern,A. and Wright,D., Weiss,R.
                                                                                                                                                                                                                                                                          Class: plasmid ends
                                                                                                                                                                                                                                                                                              Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                               Fax:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mammalia; Eutheria;
1 (bases 1 to 20)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                      308,
                                                                                                                                                                                                                                  quality sequence stop: 20.
Location/Qualifiers
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801 585 7177
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ilarity 100.0%; Pred. No.
Conservative 0; Mismatch
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/lab host="E.coli SOLER"
/lab host="E.coli SOLER"
/clone_lib="AtJMT-overexpressing transgenic rice lambda /clone_lib="AtJMT-overexpressing transgenic rice lambda phage cDNA library (JMT1)"
/note="Vector: pBluescript SK(+); Site 1: EcoRI; Site 2: XhOI; cDNA was inserted into lamda Uni-ZAP XR vector at end with EcoRI and 3' end with XhOI site. mRNA was prepared from Arabidopsis Jasmoinate Carboxyl
                /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /Clone_lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: PWA27v; Furified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jacksor
                                                                                                                                                                                                                                                                                                                                                                                                                                      Biomedical Polymers Research
Laboratory Mouse DNA Resource
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                                                                                                                                                      /organism="Mus musculus"
/mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /db_xref="taxon:39947"
/clone="JMT1--03-A01"
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/mol type="mpwa"
                                                                                                                   /db_xref="taxon:10090"
/clone="UUGC1M0009G15"
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/cultivar="Nackdong"
                                                                                                   sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chordata;
Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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                    from the Jackson
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RESULT 47
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                                                                                                                                                                                                                                                                                                                                                                                                                              Email: ddurn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0026 row: C column: 16
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1M0026C16R Mouse 10kb plasmid ŪUGC1M library Mu clone UUGC1M0026C16 R, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (2000)
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Similarity 100.0%; Pred. No. ...
O; Mismatches
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quality sequence stop: 20.
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801 585 7177
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                     /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PWD42nv, Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                        organism="Mus musculus"
/mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                  db_xref="taxon:10090"
clone="UUGC1M0026C16"
                                                                                                                                                                                                                                                                                                                                                ocation/Qualifiers
                                                                                                                                             sex="Male"
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Mouse DNA Resource
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ACCESSION VERSION KEYWORDS

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AZ366451
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Nouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                            Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0115 row: N column: 07
Seq primer: CACACAGGAAACAGCTATGACC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert B.
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/lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/clome_lib="Mouse 10kb plasmid UUGCIM library"/note="Vector: PWD42nv, Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackso: Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Biomedical Polymers Research Bldg., 20
                                                                                                                                                                                                             /mol_type="genomic DNA"
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100.0%; Pred. No. 6.
Live 0; Mismatches
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The edaptored DNA was purified and size-selected for a 9.5 to

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AZ436192
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                                                                                                                                                                                                                                                                                                                           Seq primer: CACACAGGAAACAGCTATGACC Class: plasmid ends High quality sequence stop: 20. Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0233 row: K column: 14
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Fax: 801 585 7177
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Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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/lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/clone lib="Mouse 10kb plasmid UUGC1M library"/note="Vector: PMD42nv, Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
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                                                                                                                                                                                       /mol_type="genomic DNA"
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/db_xref="taxon:10090"
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mol_type="genomic מאי"
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Insert Length: 10000 Std Error:
Plate: 0326 row: L column: 05
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University of Utah Genome Center
University of Utah
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Fax: 801 585 7177
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                                    quality sequence stop: 20.
Location/Qualifiers
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/lab host="E. Coli strain XL10-Gold, T1-resistant, F-"/Clone lib="Mouse 10kb plasmid UUGCIM library"/note="Vector: pWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Biomedical Polymers Research Bldg., 20 S. 2030
                                                                                                                               /db_xref="taxon:10090"
/clone="UUGC1M0326L05"
                                                                                                                                                                                   /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                   organism="Mus musculus"
                                                                                                            'sex≕"Male"
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100.0%; Pred. No. 6.5e+0
tive 0; Mismatches
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GSS 09-JUL-2004 i library of P.

Length 20;

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Local Similarity
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AL Submitted (23-OCT-2003) Balzergue S., UMRGV, INRA/CNRS, 2 rue Gaston Cremieux, 91057 Evry cedex, FRANCE
PCR was performed on DNA from transformants of Arabidopsis thaliana plants from INRA (Versailles). The DNA fragment(s) resulting from the PCR were directly sequenced from the left or the right border to determine the genomic sequence flanking the insertion. T-DNA derived sequences were removed. Information to order the corresponding mutant line and a link to a database providing a graphical display of the insertion site are available at http://dbsgap.versailles.inra.fr/publiclines/. This sequence has been generated in the framework of the French plant genomics program 'Genoplante' (http://www.genoplante.com and http://genoplante-info.infobiogen.fr).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Brunaud, V., Balzergue, S., Dubreucq, B., Aubourg, S., Samson, F., Chauvin, S., Bechtold, N., Cruaud, C., DeRose, R., Pelletier, G., Lepiniec, L., Caboche, M., and Lecharny, A.
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Arabidopsis thaliana (thale cress)
Arabidopsis thaliana
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EMBO Rep. 3 (12), 1152-1157 (2002)
22363535
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AJ591197.1 GI:37940821
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Arabidopsis thaliana T-DNA flanking sequence,
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                                /organism="Arabidopsis thaliana"
/mol type="genomic DNA"
/cultivar="Wassillewskija"
/db xref="taxon:3702"
/clone="581B09"
         /clone_lib="Arabidopsis thaliana T-DNA insertion lines"
                                                                                                                                                               ocation/Qualifiers
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100.0%; Pred. No. 6.5e+06;
. Mismatches 0;
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Sus scrofa (
Sus scrofa
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AJ666336
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This library was generated at Caltech,
sequenced at Vancouver, Canada.
Seq primer: T7
Class: fosmid ends.
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10; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Srinivasan,J., Otto,G.W., Kahlow,U., Geisler,R. and Sommer,R.J. AppaDB: an AcedB database for the nematode satellite organism Pristionchus pacificus
Nucleic Acids Res. 32 (1), D421-D422 (2004)
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1 (bases 1 to 20)
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/note="Vector:
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/clone_lib="Mixed stage fosmid library of P. pacificus
var. California"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Pristionchus
/mol_type="genomic DNA"
/strain="California"
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left border"
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v 100.0%; Pro
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100.0%; Pred. No.
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clone C0000033
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Pasadena, USA and end

Length 20;

Indels

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Gaps

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linear

EST 28-JUN-2004 _N17, mRNA

JOURNAL MEDLINE PUBMED REFERENCE AUTHORS

COMMENT

TITLE JOURNAL

FEATURES

source

REFERENCE AUTHORS

TITLE

ACCESSION VERSION

DEFINITION

581B09

KEYWORDS

ORGANISM

RESULT 51 AJ591197

B 5 ORIGIN

Matches Query Match Best Local

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Roslin, Midlothian, EH25 9PS, UNITED KINGDOM

Single pass sequencing. Bases called and trimmed with phred
v0.020425.c. Vector identified by cross match with the -minscore 20
and -minmatch 12 options. Vector:pBlueScriptII(KS+) R. Site 1:
ECORI R. Site 2: NotI Description: Normalised library constructed
from pooled tissue from day 30 placentas. Clones available from UK
Centre for Functional Genomics in Farm Animals, Roslin Institite,
Roslin, Midlothian, UK, EH25 9PS, www.arkgenomics.org.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    _
                                                                                                                                                                                                       Contact: Kousaku Okubo, Naohiro Hori, Ryo Matoba, Toshiyuki
Niiyama, Atsushi Fukushima, Yuko Kojima & Kenichi Matsubara
Institute for Molecular and Cellular Biology
                                                                                                                                                                                                                                                                                                                 Large scale cDNA sequencing for analysis qualitative aspects of gene expression NAT. Genet. 2, 173-179 (1992)
                                                                                                                                                                                                                                                                                                                                                                                      1 (bases 1 to 21)
Okubo, K., Hori, N., Matoba, R., Niiyama, T., Fukushima, A., Kojima, Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        D11772 21 bp mRNA linear EST 02-DEC-199 HUMHM01E05 Liver HepG2 cell line. Homo sapiens cDNA clone hm01e05,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Anderson SI
Genomics and Bioinformatics
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                    Osaka University
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                                                                                                                                       Yamada-oka, Suita, Osaka 565, Japan
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
/organism="Homo sapiens"
/mol type="mRNA"
/db_xref="GDB:D058318E"
/db_xref="taxon:9606"
/clone="hm01e05"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /db_xref="taxon:9823"
/clone="C0000033_N17"
/tissue type="placenta"
/clone_Tib="CSEQRAN09"
/clone_Tib="CSEQRAN09"
/note="Vector: pBlueScriptII(KS+); Site_1: EcoRI; Site_2:
NotI; Single pass sequencing. Normalised library
constructed from pooled tissue from day 30 placentas."
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/mol_type="mRNA"
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Class: plasmid ends
High quality sequence stop: 21.
Location/Qualifiers
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Fax: 801 585 7177
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erry
Plate: 0174 row: P column: 01
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Contact: Robert B.
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USA
was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pMD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Mus musculus"
/mol type="genomic DNA"
/strain="C57BL/6J"
/db xref="taxon:10090"
/clone="UUGC1M0174P08"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /clone_lib="Mouse_10kb_plasmid_UUGCIM_library" /note="Vector: PW42nv; Purified_genomic_DNA_from M. musculus_C57BL/6J (male) was obtained from the Jackson Laboratory Mouse_DNA_Resource_(http://www.jax.org/resources/documents/dnares/). The DN (http://www.jax.org/resources/documents/dnares/).
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/note="3'-directed regional cDNA library. Cleaved by MboI
and transformed into E.coli."
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100.0%; Pred. No. 6.5e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: ddunn@genetics.utah.edu
Insert Length: 1000 Std Error:
Plate: 0378 row: K column: 06
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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GSS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (2000)
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Location/Qualifiers
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801 585 7177
/lab host="E. Coli strain XL10-Gold, Tl-resistant, F-"
/clone_lib="Mouse_10kb plasmid_UUGCIM_library"
/note="Vector: PMD42nv; Purified_genomic_DNA_from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse_DNA Resources
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pMsm2 (gil4732114|gb|AF129072.1), a copy-number
inducible_derivative_of plasmid_R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
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/clone="UUGC1M0378K06"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /sex="Male"
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Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: C
Plate: 0429 row: H column: 03
Plate: 0429 row: ACACAGGAAACAGCTATGACC
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Mouse whole genome scaffolding with paired end reads from 10kb
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1M0429H03R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0429H03 R, genomic survey sequence.
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pMB42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mol_type="genomic DNA"
/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               clone="UUGC1M0429H03"
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Insert Length: 10000 Std Error: (
Plate: 0534 row: G column: 12
Seq primer: CACACAGGAAACAGCTATGACC
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1M0534G12R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0534G12 R, genomic survey sequence.
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Location/Qualifiers
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Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AZ658074.1 GI:11795220
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .ass: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          308,
/lab host="B. Coli strain XL10-Gold, T1-resistant, F-"
/clone lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: pWD42nv; Purified genomic DNA from M.
musculus C57BL/GJ (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.05 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWm92 (gi|4732114|gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Mus musculus"
/mol type="genomic DNA"
/strain="C57BL/GJ"
/db xref="taxon:10090"
/clone="UUGC1M0534G12"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                          sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
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Pred. No.
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COMMENT

FEATURES

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259 GATTAACAAG 268
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Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI), Shiraki
Toshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 (bases 1 to 22)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
Plate: LLAM10037 row: f column: 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       High quality sequence stop: 22.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BF979698.2 GI:12388195
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                                                                          Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
CGCCGCCGCC 48
                                                    Conservative
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                                                                                                                                                                           /clone_lib="NIH_MGC_97"
/clone_lib="NIH_MGC_97"
/clone="Organ: testis; Vector: pBluescriptR (modified /note="Organ: testis; Vector: pBluescriptR (modified /note="Organ; testis; Vector: pBluescriptR (modified pBluescript K8+); Site_1: BamHI; Site_2: SalI-XhoI
(gtcgag); Oligo-dT primed using primer
5'-TTTTTTTTTTTTTTTTTVN-3', size-selected for average insert size 2.2 kb and normalized to ROT 5. This is a primary library enriched for full-length clones and constructed using the Cap-trapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH_MGC Library."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:4374157"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                organism="Homo sapiens"
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                                            100.0%; Pr
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                                            Score 10; DB 2; Len
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; cmatches 0;
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                                                                                                Length 22;
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                                              Gaps
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RESULT 61
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Best Local Similarity
            JOURNAL
                                                                                                                                                       AUTHORS
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                                                                                                                                                                                                                                                                                                 AZ788615 22 bp DNA line 2M0035F19R Mouse 10kb plasmid UUGCIM library Mu clone UUGC2M0035F19 R, genomic survey sequence. AZ788615 GI:12928595 GSS.
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                                                                      Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                    Mus musculus
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Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Spermatoideae; Oryzeae; Oryza.
                                                  Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University YongJin, Kyeenggi, Korea Tel: 82 31 330 6193 Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Large-scale Sequencing Analysis 
Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., I
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EST
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CF309993.1 GI:33681754
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                                                                                                                                                                                                                                                                        musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note="Vector: pc44-TOPO; Site_1: EcoRI; Leaf was dried for 2hrs. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was prepared from ABA-responsive element binding transcription factor 3 overexpression line."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone_lib="ABF3-overexpressing transgenic rice plasmid cDNA library (ABF)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Oryza sativa (japonica cultivar-group)"
|mol type="mRNA"
|cultivar="Nackdong"
|db_xref="taxon:39947"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /dev_stage="14 days after germination"
/lab_host="E.coli DH10B"
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100.0%; Pred. No. 6.5e+06;
Mismatches 0;
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                 Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R. Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
                                                                                                                                                              Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia; 1 (bases 1 to 22)
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                                                                                                                                                                                                                                                                                                                                                         AZ812224 22 bp DNA linear GSS 20-FEB-200200078G13R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC2M0078G13 R, genomic survey sequence.
                                                                                                                                                                                                                                          Mus musculus
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AZ812224.1 GI:12981261
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High quality sequence stop: 22.
Location/Qualifiers
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Insert Length: 10000 Std Stror: (
Plate: 0035 row: F column: 19
Seq primer: CACACAGGAAACAGCTATGACC
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Fax: 801 585 7177
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a.
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polymucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pNB042 (gi|4732114|gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coli XL10-Gold (Stratagene) cells
and selected for ampicillin resistance."
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/clone lib="Mouse 10kb plasmid UUGC1M library"
/note="Vector: pWD42TDv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
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/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC2M0035F19"
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ive 0; Mismatches
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GSS 20-FEB-2001

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Insert Length: 10000 Std Error: (
Plate: 0078 row: G column: 13
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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Fax: 801 585 7177
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E.,
                                                             1 (bases 1 to 22)
Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R.,
Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L.,
Melville,S.E., Rajandream,M.A. and Barrell,B.G.
Direct Submission
                                                                                                                                                                                                          Trypanosoma brucei
Trypanosoma brucei
                                                                                                                                                                                                                                                                                                            genomic survey sequence.
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                   Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton,
                                                                                                                                                                                                                                                                        AL462607.1 GI:11832412
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                                                                                                                                                                                     Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
project, Sanger Centre, The Wellcome Trust Genome Car
Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                  Trypanosoma.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gil 4732114 |gb| AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/strain="C57BL/6J"
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85 GGAGGGCGAG 94
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4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. In Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
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7LEAF--02-K05.gl Rice leaf plasmid cDNA library II (7LEAF) Oryz-
sativa (japonica cultivar-group) cDNA clone 7LEAF--02-K05, mRNA
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Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; &
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Barrell, Oxford University Press, 1999).
Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger Centre are available
at http://www.sanger.ac.uk/Projects/T_brucei/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., L. Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H. Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Nahm B.H.
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82 31 321 6355
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     , Kyeonggi, Korea
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/mol_type="genomic DNA"
/strain="TREU927"
/db xref="taxon:5691"
/clone="114f05"
/dev_stage="7 days after germination"
/lab_host="8.coli DH108"
/clome_lib="Rice leaf plasmid cDNA library II (7LEAF)"
/clome_lib="Rice leaf plasmid cDNA library II (7LEAF)"
/note="Vector: pCR4-TOPO; Site 1: EcoRI; mRNA was capp:
with oligoribonucleotides and then used as templates for
                                                                                                                                            /db_xref="taxon:39947"
/clone="7LEAF--02-K05"
/tissue_type="leaf"
                                                                                                                                                                                                                                                          /organism="Oryza sativa
/mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
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                                                                                                                                                                                                                               cultivar="Nackdong"
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100.0%; Pr
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                            bhnahm@bio.myongji.ac.kr
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                                                                                                                                                                                                                                                                                           (japonica cultivar-group)"
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ORIGIN
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ORGANISM
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AZ324328/c
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.

Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       High quality sequence stop: 23.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0046 row: B column: 16
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1M0046B16F Mouse 10kb plasmid UUGC1M library Muclone UUGC1M0046B16 F, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (2000)
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801 585 7177
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                                  10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwb42 (gil 4732.14) gbl AF12907.21), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells
                                                                                                                                                                                                                                            (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"/clone lib="Mouse 10kb plasmid UUGCIM library"/note="Vector: PWD42nv; Purified genomic DNA from M.musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        clone="UUGC1M0046B16"
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100.0%; Pred. No. 6.5e+06;
... Mimmatches 0;
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               for ampicillin
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UUGC1M library Mus musculus
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JOURNAL COMMENT
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AZ345542/c
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Nouse whole genome scaffolding with paired end reads from 10kb Nouse whole genome scaffolding with paired end reads from 10kb
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University of Utah Genome Center
University of Utah
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0080 row: A column: 14
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tel: 801 585 5606
Fax: 801 585 7177
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Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1 (bases 1 to 23)
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23 bp DNA linear GSS 29-SEP-20
1M0080A14F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
2000 EUGC1M0080A14 F, genomic survey sequence.
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gh quality sequence stop: 23.
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Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was sligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into Chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /lab_host="E. Coli strain XL10-Gold, T1-resistant, P-"
/clone_lib="Mouse_10kb_plasmid_UUGC1M_library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            'clone="UUGC1M0080A14"
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100.0%; Pred. No. 6.9
/ative 0; Mismatches
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. 6.5e+06;
ches 0; Indels
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AZ761953/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0556 row: K column: 06
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ende
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R. Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AZ761953 23 bp DNA linear GSS 16-FEB-20 1M0556K06R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0556K06 R, genomic survey sequence.
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Location/Qualifiers
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Fax: 801 585 7177
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             /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWM92 (gil4732114|gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coli XL10-Gold (Stratagene) cells
and selected for ampicillin resistance."
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/clone="UUGC1M0556K06"
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/strain="C57BL/6J"
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University of Utah
University of Utah
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Nouse whole genome scaffolding with paired end reads from 10kb Mouse whole genome scaffolding with paired end reads from 10kb
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Insert Length: 10000 Std Error:
Plate: 0216 row: C column: 19
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of phw92 (gi[473114]gb[AF19072.1], a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/clone_lib="Mouse 10kb plasmid UUGC2M library"
/note="Vector: pWD42nv, Purified genomic DNA from M.
musculus_C57BL/6J (female) was obtained from the Jackson
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/clone="UUGC2M0216C19"
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/strain="C57BL/6J"
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Matches 10
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AG198216.1
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1 Similarity 100.0%;
10; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger Centre
at http://www.sanger.ac.uk/Projects/T_brucei/.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 to give a tight size distribution (4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Pan.
                                                                                   Pan
                                                                                                                                GSS.
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                                                                                                       Pan
                                                                                                                                                                                                                              Pan troglodytes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned population of Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically sheared
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          T. brucei sheared genomic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Trypanosoma brucei
Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
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/clone="125f10"
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/strain="TREU927"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        organism="Trypanosoma brucei"
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100.0%; Pred. No. 6.5e+06;
tive 0; Mismatches 0;
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RP43-078005.TJ, genomic
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o. 6.5e+06;
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                                Unpublished
2 (bases 1 to 23)
2 (bases 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           272 AAGGTTGTTA 281
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Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequence.
AG202522
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Clones are derived from the chimpanzee BAC library RP-43 This BAC end was generated during the R&D process and may have higher chance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pan troglodytes (chimpanzee)
Pan troglodytes
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Park, H., Kim, Y., Kim, S., Han, Y., Woo, T., Pathoon, S.T., Chu, M., Kim, H., Joo, S., Kim, C.,

Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Pan.
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LIBRARY
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Sequencing: TJ
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R.Site 1 : EcoRI
R.Site 2 : EcoRI.
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/db_xref="taxon:9598"
/clone="RP43-078005.TJ"
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'clone_lib="RP-43 Chimpanzee Male BAC Library"
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0; Mismatches 0;
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V. and Yoo, H.
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RESULT 73
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tetrahymena thermophila
Eukaryota; Alveolata; Ciliophora; Oligohymenophorea;
Hymenostomatida; Tetrahymenina; Tetrahymena.
    AZ362057
                                                                                                                                  h 3.1%; Sco
Similarity 100.0%; P
                                                                                                                                                                                                                                                                                                                                                                                                                       Molecular Genetics and Cell Biology
University of Chicago
920 E. 58th Street, Chicago, IL 60637, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EST from Tetrahymena thermophila, strain CU428.1, growing cells Unpublished (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Turkewitz, A.P., Karrer, K.M., Frankel, J. and Klobutcher, L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BM400127 24 bp mRNA linear EST 17-JAN-2002 5009-0-66-G02.t.2 Chilcoat/Turkewitz cDNA (large fraction)
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                                                                                                                                                                                                                                                                                                                                                                Seq primer: T3
                                                                                                                                                                                                                                                                                                                                                                             Email: apturkew@midway.uchicago.edu
                                                                                                                                                                                                                                                                                                                                                                                              Tel: 773 702 4374
Fax: 773 702 3172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Turkewitz AP
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                                                                                                                                                                                                           /db_xref="taxon:5911"
/clone lib="Chilcoat/Turkewitz cDNA (large fraction)"
/note="Vector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
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ite 1 : EcoRI
ite 2 : EcoRI.
Location/Qualifiers
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/mol type="genomic DNA"
/db_xref="teaxon:9598"
/clone="RP43-086A14.TJ"
/sex="male"
                                                                                                                                                                                                                                                                                      organism="Tetrahymena thermophila"

mol_type="mRNA"

strain="CU428.1"
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/clone_lib="RP-43 Chimpanzee Male BAC Library"
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                                                                                                                                                 Score 10; DB 4; L
Pred. No. 6.5e+06;
                                                                                                                                      Mismatches
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DEFINITION
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Best Local Similarity
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Insert Length: 1000 Std Error: 0.00
Plate: 0107 row: E column: 05
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
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A2362057
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Rese
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801 585 7177
      Conservative
                                                                                                                                          /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWM92 (gil4732114|gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coli XL10-Gold (Stratagene) cells
and selected for ampicillin resistance."
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/mol_type="genomic DNA"
/strain="C57BL/6J"
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3.1%; Score 10; DB 8; Le
100.0%; Pred. No. 6.5e+06;
tive 0; Mismatches 0;
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University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E.,
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Insert Length: 10000 Std Error: 0.00
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801 585 7177
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                                                                                                                                                                                                                                                                                                                                 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          source
232 GGAAGATACC 241
                                               10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Class: plasmid ends
Class: plasmid ends
High quality sequence stop: 24.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R. Mouse whole genome scaffolding with paired end reads from 10kb
                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
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Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 24)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mus musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AZ648378.1 GI:11780785
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                                                     Conservative
                                                                                                                                                                    (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pW42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"/Clone lib="Mouse 10kb plasmid UUGCIM library"/note="Vector: pW042nv; Purified genomic DNA from M.musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /db_xref="taxon:10090"
/clone="UUGC1M0517D24"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mol_type="genomic DNA"
strain="C57BL/6J"
                                                                            100.0%;
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Pred. No.
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1 20 6.3 28 5 AAF30355 2 2 16 5.0 20 2 AAV09174 3 15 4.7 18 2 AAX63292 4 15 4.7 18 2 AAX63292 A AAX63292 A AAX63294 C 5 15 4.7 25 9 ACI37850 A ACI37850 AACI37850 AACI37850 AACI37850 AACI37850 A ACI37850	2: geneseqn1980s:* 3: geneseqn2000s:* 4: geneseqn2001bs:* 5: geneseqn2001bs:* 6: geneseqn2001bs:* 7: geneseqn2003bs:* 9: geneseqn2003bs:* 10: geneseqn2003bs:* 11: geneseqn2003ds:* 12: geneseqn2003ds:* 12: geneseqn2004ds:* 12: geneseqn2004s:* 13: geneseqn2004s:* 14: geneseqn2004s:* 15: geneseqn2004s:* 16: geneseqn2004s:* 17: geneseqn2004s:* 18: geneseqn2004s:* 19: geneseqn2004s:* 19: geneseqn2004s:* 10: geneseqn2004s:* 11: geneseqn2004s:* 12: geneseqn2004s:* 13: geneseqn2004s:* 14: geneseqn2004s:* 15: geneseqn2003cs:* 16: geneseqn2003cs:* 17: geneseqn2003bs:* 18: geneseqn2003bs:* 19: geneseqn2003bs:* 10: geneseqn2003cs:* 10: geneseqn2003cs:* 10: geneseqn2003cs:* 11: geneseqn2003cs:* 12: geneseqn2003cs:* 13: geneseqn2003bs:* 14: geneseqn2003bs:* 15: geneseqn2003cs:* 16: geneseqn2003cs:* 17: geneseqn2003cs:* 18: geneseqn2003cs:* 19: geneseqn2003cs:* 10: geneseqn2003cs:* 10: geneseqn2003cs:* 10: geneseqn2003cs:* 10: geneseqn2003cs:* 11: geneseqn2003cs:* 12: geneseqn2003cs:* 13: geneseqn2003cs:* 14: geneseqn2003cs:* 15: geneseqn2003cs:* 16: geneseqn2003cs:* 17: geneseqn2003cs:* 18: geneseqn2003cs:* 19: geneseqn2003cs:* 10: geneseqn200as:* 10: geneseqn200as:* 10: geneseqn200as:* 10: geneseqn20	DB seq leng cocessing: Li	4134886 : 0 er of hits sa	Title: US-10-048-046-1_COPY_81_399 Perfect score: 319 Sequence: 1 gtgaatcccgatggagcggctacagactggggatgtcatc 319 Scoring table: OLIGO_NUC Gapop_60.0 , Gapext 60.0	GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd. OM nucleic - nucleic search, using sw model Run on: February 2, 2005, 15:06:25; Search time 149.916 Seconds (without alignments) 11170.029 Million cell updates/sec
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ABV89390 ABV89390 ABV89389 ACA09011 ACA09012 ACA09012 ACA09012 ACA09012 ACA09013 ACA090366	ABC96919 ABF95242 ABF95243 ABF95243 ABC96918 AAA2613 ABN262530 ABN26230 ABN00430 ABN00430 ABN00430 ABN0046453 ABN131182 ABV78966	ACK00130 2 ADP15727 0 ABQ80298 2 ADK00061 ABX67739	2 ADO09916 AA141635 0 ADH93941 2 ADK98335 AAQ34273 ACK00756 ACI86610	0 ADF51980 0 ADF51980 2 ADP05478 2 AAV09175 AAX93253 AAC67142 ABC67142 ABC67142 ABC67142	0 ADF52686 0 ADF52703 0 ADF52726 0 ADF5293 0 ADF52685 0 ADF52685 0 ADF52686 0 ADF52698 0 ADF52698 0 ADF52007 0 ADF52007
Abv89392 Human PÓS Abv89390 Human PÓS Abv89399 Human PÓS Abv89389 Human PÓS Aca09011 NFKB sub- Aca09011 NFKB sub- Aca09012 NFKB sub- Acd57027 HCV DNAZY Adi82871 HCV DNAZY Adv61703 Hepartitis Aav30545 HLA DQB1 Abl30595 Human HLA Abl30696 Human HLA Abl30696 Human HLA Add69478 3' anchor Abk29933 Cyclin D1 Ach03515 Human lat Add00366 HCV codin Adf35778 Human VEG Adf36705 Human VEG Adf52709 Hepatitis	AGE89508 HIV-1 REV ADC96919 Oligonucl Ab695242 Oligonucl Ab695243 Oligonucl Abc96918 Oligonucl Abc96918 Oligonucl Aba26133 Oestrogen Aba26133 Oestrogen Aba262579 Substrate Abx00430 Hepatitis Aa262579 Substrate Abx00430 Hepatitis Aa26476 Probe DB6 Aaq26453 Probe DB6 Aaq26453 Probe DB6 Abl31182 Human HTP Abv78964 Human HTP Abv78965 Human HTP Abv78965 Human HTP Abv78966 Human HTP Abv78966 Human HTP Abv78966 Human ERG Abk19313 Human ERG Abk19315 Human ERG Abk19315 Human POS Abv89388 Human POS	Ack00130 Human mic Ack00130 Human mic Adp15727 Renal cel Abg80298 Primer Te Adk00061 Human tel Abx67739 Novel Hel	Ado09916 Human NOV Aa141635 Human col Adh93941 Human gen Adk98335 Primer of Aag4273 Upstream Ack00756 Human mic Aci86610 Human mic	Adf51969 Hepatitis Adg05478 Hepatitis Adg05478 Hepatitis Aav09175 Phosphoro Aax93253 PCR prime Aac67142 Human E2F Abt0851 Human nov Adj34405 Human sec	Adf52686 Hepatitis Adf52703 Hepatitis Adf52726 Hepatitis Adf52726 Hepatitis Adf52030 Hepatitis Adf52994 Hepatitis Adf52685 Hepatitis Adf52689 Hepatitis Adf52697 Hepatitis Adf52990 Hepatitis Adf52990 Hepatitis Adf52690 Hepatitis Adf52092 Hepatitis

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Checkpoint with forkhead associated domain and ring finger; Chfr; human; mitosis; cell cycle; tumour; diagnosis; antitumour; drug screening;
                                                           Human checkpoint gene chfr 5' PCR primer.
                                                                                            14-MAY-2001
                                                                                                                                                           AAF30355 standard; DNA;
                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                        ADG29865

ADG2145

AAG37582

AAT64896

AB182437

AB182437

AB192632

AB192632

AB192632

AB192702

AB191702

AB191703

AB19170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAZ35710
AAF87680
ABN86474
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ADF86274
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ADM48621
ADN48691
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ADP56863
AAT89731
AAV47454
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ADF52013
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AAQ21051
AAT89745
AAZ07024
ABL42978
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ADF37724
ADG29869
ADJ97418
AAA28540
ACA64608
AAQ37589
AAT64903
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ABT15791
ABT43241
ABT32412
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ADF37816
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Aat64896 Hepatitis
Abi82437 Capture o
Abi82436 Capture o
Abi92632 Capture o
Abi92832 Capture o
Abi92833 Capture o
Abi92702 Capture o
Abi92833 Capture o
Abi91703 Capture o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Abt15791 Human GU
Abt13741 Neuroblas
Abt32412 Neuroblas
Abt32412 Neuroblas
Abt32412 Neuroblas
Acd23026 Human NEM
Adf86274 Rat GAPDH
Add78398 Human E2-
Adn48621 Human Not
Adn56791 Artisense
Adn556791 Artisense
Adp56791 Artisense
Aat89721 PCR prime
Aat89721 PCR prime
Aat89731 Human uro
Aaf87680 Human MP
Add14547 Human WP
Add14547 Human VEG
Add29869 PLT1-targ
Add397418 Human Flt
Aaa28540 PCR primer
Aca64608 C. elegan
Aad37589 HCV conse
Aat64903 Hepatitis
Abk66685 Human VEG
Add37816 Human VEG
Add37818 FLT1-targ
Ado22145 Real-time
Aaq37982 HCV conse
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Aat89745 Oligonucl
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Adf52013 Hepatitis
Adf52452 Hepatitis
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6 HLA DQB g
6 HLA DQB1
6 M.f.cytb2
7 Human HTP
6 Human HTP
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Human chr
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          RESULT 2
AAV09174/c
ID AAV09174 standard;
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Matches 20
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                                                Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel nucleic acid sequence of mitotic checkpoint gene encoding a checkpoint with forkhead-associated domain and ring finger protein, diagnosing tumorigenic cells and in screening for anticancer drugs.
                                                                                                        Homo
                                                                                                                       Synthetic.
                                                                                                                                                  Type I interleukin-1 inflammation; ss; 5'
                                                                                                                                                                                                  Phosphorothiate oligonucleotide sequence
                                                                                                                                                                                                                                     09-JUN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 28
                                                                                                                                                                                                                                                                   AAV09174;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 3; Page 38; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-182927/18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Halazonetis T, Scolnick D;
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                                                                                                      sapiens
                                                                                                                                                                                                                                                                                                                                                                                                    230 CTGGAAGATACCAGCACCAG 249
                                                                                                                                                                                                                                                                                                                                                                                                                                                20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 12 A; 6 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                  (first entry)
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                                                      Location/Qualifiers
1. .20
                                     /*tag=
                                                                                                                                                                                                                                                                                                  DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               6.3%;
100.0%;
                                                                                                                                                  receptor; IL1R; human; IL1 protein; hybridisation; Cap region; phosphorothioate linkage.
        a
"Phosphorothioate internucleotide linkage"
                                                                                                                                                                                                                                                                                                    20
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Mismatches . 6;

0; Indels Length 28;

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8051 targeting IL1R mRNA

RESULT 1
AAF30355
ID AAF3
XX
AC AAF3
XX
DT 14-M
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DE Huma
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KW Chec
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105 106 107 108 109 110 111 111 111

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RESULT 3
AAX63292
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XX AAX6
AC AAX6
AC AAX6
XX Delt
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KW Maiz
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                 Zwick MG,
Young SA,
                                                                                                                                                                                                                                                                                                                                                              Maize; corn; Zea maye; delta-9 desaturase; GBSS; target; substrate; granule bound starch synthase; hammerhead ribozyme; hairpin ribozyme; modulation; gene expression; transgenic plant; cleavage; canola plant caffeine synthesis; coffee plant; nicotine production; tobacco;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-JUL-1999
                                                                                                                                                                                                                                                                                                        Zea mays.
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type I - used to modulate expressi
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                                                                                                                                                                                                                           20-MAR-1997
                                                                                                                                                                                                                                                                    WO9710328-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Delta-9 desaturase hairpin ribozyme substrate SEQ ID NO:1167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAX63292 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
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                                                                               (RIBO-)
                                                                                                                                         13-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-MAY-1996;
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                                                                                                                                                                                  12-JUL-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               therapeutic purposes
                                                                                                                                                                                                                                                                                                                                                 ripening; flower pigmentation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19
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                                                                             RIBOZYME PHARM DOWELANCO.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GCCGCCGCAGCCCTGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                         95US-0001135P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RNA;
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100.0%; Pr
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               Mcswiggen
Merlo DJ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     otide(s) specific for interleukin-1 receptor expression and detect overexpression of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dean
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Pred.
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7e+02;
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RESULT 4
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Best Local
                                                                                                                  Zwick MG,
Young SA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention describes an enzymatic nucleic acid molecule (I) with RNA cleaving activity, which modulates the expression of a plant gene. Also described is a gene comprising a cDNA sequence encoding maize Delta-9 desaturase. (I) can be used to modulate expression of a gene, preferably Delta-9 desaturase or a granule bound starch synthase (GBSS) gene, in a plant (preferably a maize or canola plant). (I) can be used to modulate caffeine synthesis in a coffee plant, nicotine production in a tobacco plant, fruit ripening processes in an apple, tomato, pear, plum or peach plant, flower pigmentation in a rose, petunia, chrysanthemum or marigold plant or lignin production in a tobacco, aspen, poplar or pine
                                            Ribozyme which modulates plant gene expression of DELTA-9 desaturase or maize or canola.
                                                                                                                                                                                                                                                                                                                           granule bound starch synthase; hammerhead ribozyme; hairpin modulation; gene expression; transgenic plant; cleavage; carfeine synthesis; coffee plant; nicotine production; tobac
                                                                                                                                                       (RIBO-)
                                                                                                                                                                                           13-JUL-1995;
                                                                                                                                                                                                                  12-JUL-1996;
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                                                                                                                                                                                                                                                                                                                                                                                      Delta-9 desaturase hairpin ribozyme substrate SEQ ID NO:1169.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX63294 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     maize or canola.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ribozyme which modulates plant gene expression - preferably modulates expression of DELTA-9 desaturase or granule bound starch synthase in
                                                                                            WPI; 1997-202224/18
                                                                                                                                                                                                                                                                                          Zea mays.
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                                                                                                                                                                                                                                                                                                               ripening;
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Folkerts
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                                                                                                                                                                                                                                                                                                                                                               Zea mays; delta-9 desaturase;
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                                                                                                                                                                                                                                                                                                               flower pigmentation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               RNA;
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Merlo DJ;
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                                                                                                                               JA,
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J. 2.3e+03;
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                                                                                                                                PAO,
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                                                        preferably modulates
d starch synthase in
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                                                                                                                                                                                                                                                                                                                                       canola plant;
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The present invention describes an enzymatic nucleic

acid molecule

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Claim 40;

93; 155pp;

English.

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The invention discloses a microarray comprising a plurality of nucleic cald probes including one of 2,018,500 fully defined sequences, or its perfect match, perfect mismatch, antisense match or antisense mismatch. Also disclosed is a method of gene expression analysis. The array is used in monitoring gene expression levels by hybridisation to a DNA library, in analysis of genetic variation or in hybridisation of tag-labelled compounds. The nucleic acid probes are specifically designed for analysis of fat least one target sequence. The method of analysis comprises hybridising at least one or more nucleic acids to at least two or more nucleic acids to at least two or more nucleic acids to at least one or more nucleic acids to at least or polymorphisms, or family members of a gene and a cross-species comparison. Each of the nucleic acids further comprises a tag sequence. The array of nucleic acids further comprises a tag sequence. The array of nucleic acid
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ACI37850/c
ID ACI37850 standard; DNA; 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New array of nucleic acid probes, useful for Southern, Northern or dot-blot hybridization sequence or specific mutations of any gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; SEQ ID NO 37841;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-567953/53
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-MAR-2002; 2002US-00098263.
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Pred. No.
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hybridisation,

Southern, Northern or dot-

Query Match
Best Local Similarity
Matches 15; Conserv

4.7%; ilarity 100.0%; Conservative

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Score 15; DB; Pred. No. 2.3

DB 6; Le

Length

27

Indels

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Gaps

<u>.</u>

Sequence

27

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5 A;

3 C; 10

G; 9 T; 0 U; 0 Other;

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RESULT 6
AAL42088
ID AAL4
XX AAL4
XX AAL4
XX AAL4
XX AAL4
XX Huma
XX Huma
XX Homc
OS Synt
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                                                   The invention comprises a method for detecting the presence of alleles of the glutathione S-transferase (GST) genes GSTM1, GSTM3, GSTP1 and GSTT1. The method of the invention involves the geoffically claimed GST allele-specific PCR primers. Cancer research has shown that the presence of various GST polymorphisms correlates with altered risk for certain cancers and altered response and toxicity from currently used cancer treatments (e.g. chemotherapy). The method of the invention is useful for identifying the alleles of GST present in a sample of genetic material and characterising the genetic makeup of a subject. The present DNA sequence represents the GSTT1 allele-specific PCR primer (T1R)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          blot hybridisation to identify or detect the sequence or specific mutations of any gene, in mapping the 5 termini of mRNA molecules by primer extensions or in screening cDNA or genomic libraries or subclones for additional subclones containing segments of DNA that have been isolated and previously sequenced. The sequence presented is one of the nucleic acid probes incorporated in the microarray. Note: The sequence data for this patent can also be obtained in electronic format directly from USPTO at seqdata.uspto.goc/sequence.html
                                                                                                                                                                                                                                                                                                                                                                                                     Claim 29; Page 5; 37pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; ss; allele-specific; PCR; primer; glutathione S-transferase;
GSTM1; GSTM3; GSTP1; GSTT1; cancer; cancer treatment; chemotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-241577/29.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-JUL-2000; 2000US-0219531P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-JUL-2001; 2001WO-US022923
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200208465-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human Glutathione S-transferase GSTT1 allele-specific PCR primer (T1R).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-MAY-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (UTAH ) UNIV UTAH RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   176 GTCTCTGGAGATCAC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      characterisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GTCTCTGGAGATCAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                glutathione S-transferase alleles, by amplifying portions of NA samples with primers specific for GSTM1 and CDK2, for GSTT, and for GSTP1, and detecting alleles and gene dosages.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ballard L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FOUND.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lemons R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ω
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ali-Osman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 9; L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for GSTT1
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RESULT 7
  RESULT 8
ACD65641
ID ACD6
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AC ACD6
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DT 30-5
XX
DE HCV
XX
KW Nucl
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Best Local Similarity |
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                                                                                                                                                                                                                                                                                                            This invention relates to a human purine receptor polypeptide (a P2Y receptor), termed TG20022 selected from a 331 amino acid sequence, given in the specification. The receptor of the invention is a novel G-protein coupled receptor, involved in signal transduction in many different pathways. The polypeptide is useful for screening or identifying a ligand, agonist or antagonist of the receptor of the invention, or for detecting the function or activity of the polypeptide. The polypeptide is useful as a therapeutic agent. The present sequence is that of a PCR primer which was used for amplification of a region of the human TG20022 gene during the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   purine receptor; TG20022; novel G-protein coupled receptor; signal transduction; human; P2Y receptor; PCR; primer; ss.
Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense;
                                        HCV minus
                                                                  30-SEP-2003
                                                                                                                  ACD65641 standard; RNA; 17
                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel purine receptor polypeptide, termed TG20022, useful for screening or identifying ligand, agonist or antagonist of purine receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-JUL-2001; 2001JP-00206914.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human purine receptor TG20022 PCR primer Seq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADC55392;
                                                                                           ACD65641;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-700043/67.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-JUL-2001; 2001JP-00206914.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12-FEB-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADC55392
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (TANA ) TANABE SEIYAKU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sapiens
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                                                                                                                                                                                                                                                                                       28 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TACAGACTGGGGATG
                                                                                                                                                                                                          AGGTGACACTGGAAG 236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TACAGACTGGGGATG 20
                                        strand
                                                                                                                                                                                                                                   Conservative
                                                                  (first entry)
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                                                                                                                                                                                                                                                                                       9
                                       DNAzyme substrate sequence #2160.
                                                                                                                                                                                                                                                                                       Α,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
                                                                                                                                                                                                                                                4.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           9,
                                                                                                                                                                                                                                                                                     7 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8
                                                                                                                                                                                                                                                                                                                                                                                                                                                           23pp; Japanese.
                                                                                                                                                                                   28
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                                                                                                                     ВP
                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                Score 15;
Pred. No.
                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                             DB 10;
                                                                                                                                                                                                                                              2.3e+03
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                                                                                                                                                                                                                                                          Length 28
                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                   Gaps
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닭 S

RESULT 9

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The present invention relates to nucleic acid molecules which modulate CC the synthesis, expression and/or stability of Hepatitis C virus (HCV) or CC Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, CC inozymes, zinzymes, amberzymes, and G-cleaver ribozymes. Also disclosed are nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV DNA. The nucleic acids may be used to modulate the expression of HBV CC DNA. The nucleic acids may be used to modulate the expression of HBV CC DNA. The nucleic acids may be used to modulate the expression of HBV CC DNA. The nucleic acids may be used to modulate the expression and/or replication of HCV. The compounds that modulate the expression and/or replication of HCV. The compounds and methods of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene expression such as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence represents substrate for one of the HCV converge.
                                                                                Matches
                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Blatt L, ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-JUN-2001;
08-JUN-2001;
24-OCT-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; zinzyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic;
                                                                                                                                                                 Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel compound useful for hepatocellular carcinoma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (RIBO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-MAR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  virucide; antiinflammatory; substrate; ss.
                                                                                                                                                                                                                                 DNAzyme or minus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Page 313; 387pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (MACE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-DEC-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hepatitis C virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (LEEP/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (PAVC/)
                                                                                                      OCal
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                                      139 ACGAGGTTGCGACC
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DRAPER K.
ROBERTS E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MCSWIGGEN
MORRISSEY
                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RIBOZYME PHARM
BLATT L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PAVCO P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MACEJAK
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Roberts E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; 2001US-00817879.
; 2001US-00877478.
; 2001US-0296876P.
; 2001US-0335059P.
                                                                                                                                                                   BP; 3
                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2001US-0337055P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ö
                                                                                                                                                                                                                                 strand DNAzyme sequences disclosed
                                                                                                                                                                   P
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                                                                                                4.4%;
                                                                                                                                                                 6 C; 5 G; 0 T; 3 U; 0 Other;
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Pred. No. 7.5e+03
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The present invention relates to nucleic acid molecules which modulate CC the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense include antisense since the synthesis, actives the such as hammerhead ribozymes, DNAzymes, C inozymes, zinzymes, and G-cleaver ribozymes. DNAzymes, c inozymes, zinzymes, and G-cleaver ribozymes. Also disclosed are nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV CC DNA. The nucleic acids may be used to modulate the expression of HBV CC genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and disease states related to HBV and HCV infection, replication and gene capression such as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence represents a substrate for one of the HCV invention or minus strand DNAzyme sequences disclosed in the present
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ID ACD57028 standard;
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08-JUN-2001;
24-OCT-2001;
05-DEC-2001;
                                                                                                                                                                                                                                                                                                         Claim
                                                                                                                                                                                                                                                                                                                                                 Novel compound useful for hepatocellular carcinoma,
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(BLAT/)
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LEE P.
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MORRISSEY D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      acid molecule; Hepatitis C virus; HCV; Hepatitis B virus;
                                                                                                                                                                                                                                                                                                       Page
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Roberts E;
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; 2001US-00877478.
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Matches 14
The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, inozymes, zinzymes, amberzymes, and G-cleaver ribozymes. Also disclosed are nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase and/or HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV DNA. The nucleic acids may be used to modulate the expression of HBV
                                                                                                                                                                                                                                                                                                                                                                                                                                           Blatt I
Draper
                                                                                                                                                                                                                                                    Claim 1; Page
                                                                                                                                                                                                                                                                                                                   Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis
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(BLAT/)
(MACE/)
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24-OCT-2001;
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HBV reverse transcriptase; Enhancer I region; viral replication;
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DRAPER K.
ROBERTS E.
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MORRISSEY
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BLATT L.
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Roberts E;
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; 2001US-0296876P.
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in the phenomena of tumour suppression, tumour reversion, apoptosis and/or resistance to viruses. The invention may be useful for the development of compounds with a cytostatic, virucide, neuroprotective, nootropic or neuroleptic activity. The DNA sequences may be useful as probes and primers for detecting, indentifying, quantifying and/or amplifying nucleic acid, for example as one component of a gene chip, in vitro as antisense reagents and for production of recombinant polypeptides. The invention may therefore be useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia. The present sequence is that of a nucleic acid sequence of the invention. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and methods of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene expression such as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence represents a substrate for one of the HCV DNAzyme or minus strand DNAzyme sequences disclosed in the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   tumour suppression; tumour reversion; apoptosis; virus resistance; cytostatic; virucide; neuroprotective; nootropic; neuroleptic; probe; primer; PCR; gene chip; antisense; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; ds; hu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New isolated nucleic acid, useful with tumors and cell degeneration, and transfected cells.
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                                                                        Sequence
                                                                                                                                                                             Claim 1; SEQ ID NO 4437; 198pp; English
                                                                                                                                                                                                Enzymatic nucleic acid molecules which specifically cleave RNA derived from hepatitis C virus (HCV), useful for the treatment of HCV infection especially in combination with type I interferon therapy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Blatt L,
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            This invention relates to a novel nucleic acid fragment for hybridising target nucleic acid which contains a first target sequence and a second target sequence, which comprises a sequence complementary to the first
                                                                                                                                                                     Novel nucleic acid fragment comprising sequence complementary to first target sequence, second target sequence and spacer, useful as nucleic acid probe for detecting target nucleic acid and as primer for extending target nucleic acid and as primer for extending target nucleic acid.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel nucleic acid fragment comprising sequence complementary and second target sequence and arbitrary sequences, useful as detecting hepatitis-C virus or for hybridizing target nucleic
                                                                                                                    Example 1;
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20-DEC-2001;
01-FEB-2002;
                                                                            The invention relates to a novel isolated double stranded RNA oligonucleotide about 19 to about 25 ribonucleotides in length or its equivalent. One strand of the oligonucleotide comprises the same nucleotide sequence as a region of a hepatitis C virus (HCV) target RNA polynucleotide sequence required for hepatitis C virus infection, replication or pathogenesis in vitro or in vivo in a host cell. The oligonucleotide of the invention demonstrates virucide activity and may be useful for preparing a composition or vaccine for treating or preventing hepatitis C virus, as well as during gene therapy procedures. The current sequence is that of the HCV coding region-derived conserved RNA sequence of the invention.
                                                                                                                                                                                                                                                                 New double stranded RNA oligonucleotide, useful for preparing composition for treating or preventing hepatitis {\tt C} virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention is useful as a reagent such as a nucleic acid probe for detecting target nucleic acid or as a primer for extending target nucleic acid. In particular, the invention is useful for detecting human hepatitis-C virus. The invention enables a highly sensitive and specific detection of a target nucleic acid, as a probe, and efficient extension of target nucleic acid, as a primer. The present sequence is that of an oligonuclectide probe which was used for detection of a region of the
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                                                                                                                                                                                                                                        Page 70;
  Conservative (
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01-JAN-2004 ADD00363;

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RESULT 17
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                                                                                                                                       Query Match
Best Local
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                                                                                                                                                                                                                           Sequence 19 BP; 4 A; 5 C; 7 G; 0 T; 3 U; 0 Other;
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20-DEC-2001; 2001US-0344116P
01-FEB-2002; 2002US-0353750P
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                                                                                                               Conservative
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100.0%;
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17-AUG-2001; 2001US-0313076P.
                               16-AUG-2002; 2002WO-US021843.
                                                             27-FEB-2003
                                                                                           WO2003016572-A1
                                                                                                                        Hepatitis C virus
                                                                                                                                                    gene therapy;
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20-DEC-2001; 2001US-0344116P
01-FEB-2002; 2002US-0353750P
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                                                                  New double stranded RNA oligonucleotide, useful for preparing composition for treating or preventing hepatitis C virus.
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20-DEC-2001; 2001US-0344116P
01-FEB-2002; 2002US-0353750P
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                       ASS, and where most of the pyrimidine nucleotides comprise a sugar modification. The invention may allow development of compounds with virucide, antiinflammatory, hepatotropic or cyrostatic activities by modulation (inhibition) of expression or activity of HCV RNA, by RNA interference. The siNA's of the invention may be used to inhibit replication of HCV, in cells, tissue explants or organisms, for treating HCV infection and its consequences (liver failure; hepatocellular cancer and cirrhosis), and also for drug screening, diagnosis, target identification and validation, genetic engineering, pharmacogenomics, studying gene function and gene mapping (for example of single-nucleotide polymorphisms). The chemical modification improves stability, activity, cellular uptake and/or binding affinity. The slNA can be directed to conserved regions of HCV genes, so are active against many different
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11-MAR-2002;
16-MAR-2002;
26-MAR-2002;
06-JUN-2002;
05-AUG-2002;
                                                                                                                                                                                                                                     This invention relates to novel double-stranded short interfering nucleic acids (siNA) that inhibits replication of hepatitis C virus (HCV), where one strand is an antisense strand (ASS) that is complementary to (part
                                                                                                                                                                                                                                                                                                  Example
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09-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-FEB-2003;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    short interfering nucleic acid; siNA; virus replication inhibition; hepatitis C virus; HCV; sugar modification; virucide; antiinflammatory; hepatotropic; cytostatic; RNA interference; HCV infection; liver failure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hepatitis C virus siNA antisense strand SeqID1276.
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                                                                                                                                                                                                                                                                                                                                                                                                                    Mcswiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                     (SIRN-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hepatocellular cancer; cirrhosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22
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                                                                                                                                                                                                                       strand is an antisense strand (ASS) that is complementary to (part an HCV RNA (portion) and a sense strand (SS) that is complementary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             139 ACGAGGTTGCGACC 152
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14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                     SIRNA
                                                                                                                                                                                                                                                                                                 <u>υ</u>
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                                                                                                                                                                                                                                                                                               SEQ ID NO 1276; 183pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2002US-0363124P
2002WO-US009187.
2002US-0386782P
2002US-0401104P
2002US-0406784P
2002US-0408378P
2002US-0408378P
2002US-0408378P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ilarity 100.0%; | Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2003US-0440129P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 14; DB; Pred. No. 7.5
                                                                                                                                                                                                                                                                                                                                                                                                                      Macejak D,
                                                                                                                                                                                                                                                                                                                                           g nucleic treating
                                                                                                                                                                                                                                                                                                                                                                                                                      Morrissey
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hes 0;
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                                                                                                                                                                                                                                                                                                                                            acid comprises infection with
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hepatitis
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RESULT 23
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    This invention relates to novel double-stranded short interfering nucleic cacids (siNA) that inhibits replication of hepatitis C virus (HCV), where one strand (ASS) that is complementary to (part of) an HCV RNA (portion) and a sense strand (ASS) that is complementary to ASS, and where most of the pyrimidine nucleotides comprise a sugar modification. The invention may allow development of compounds with virucide, antiinflammatory, hepatotropic or cytostatic activities by modulation (inhibition) of expression or activity of HCV RNA, by RNA interference. The siNA's of the invention may be used to inhibit creplication of HCV, in cells, tissue explants or organisms, for treating HCV infection and its consequences (liver failure, hepatocellular cancer and cirrhosis), and also for drug screening, diagnosis, target clading gene function and gene mapping (for example of single-nucleotide polymorphisms). The chemical modification improves stability, activity, consequenced reading affainty. The siNA can be directed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local
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11-MAR-2002; 2002US-0363124P.
26-MAR-2002; 2002WS-0386782P.
06-JUN-2002; 2002US-0401104P.
29-AUG-2002; 2002US-0406784P.
29-AUG-2002; 2002US-0406378P.
05-SEP-2002; 2002US-0409293P.
15-JAN-2003; 2003US-0440129P.
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                                                                                                                                                                                                                                                                                                                New double-stranded short interfering nucleic modified pyrimidine bases useful for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      short interfering nucleic acid; siNA; virus replication inhibition; hepatitis C virus; HCV; sugar modification; virucide; antiinflammatory; hepatotropic; cytostatic; RNA interference; HCV infection; liver failure;
                                                                                                                                                                                                                                                                       Example 3;
                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-689778/65
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hepatocellular cancer; cirrhosis; вв.
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12; Conserv
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                                                                                                                                                                                                                                                                    SEQ ID NO 1293; 183pp; English
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85.7%;
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                                                                                                                                                                                                                                                                                                                                                                                          Macejak D,
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                                                                                                                                                                                                                                                                                                                                                                                        Morrissey
                                                                                                                                                                                                                                                                                                              acid comprises sugar-
infection with hepatitis
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RESULT 24
ADF52726
ID ADF52
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This invention relates to novel double-stranded short interfering nucleic cacids (siNA) that inhibits replication of hepatitis C virus (HCV), where cone strand is an antisense strand (ASS) that is complementary to (part CC) an HCV RNA (portion) and a sense strand (SS) that is complementary to ASS, and where most of the pyrimidine nucleotides comprise a sugar compdification. The invention may allow development of compounds with convicte, antinflammatory, hepatotropic or cytostatic activities by compound inhibition) of expression or activity of HCV RNA, by RNA interference. The siNA s of the invention may be used to inhibit consequences (liver failure; hepatocellular cancer confidentification and its consequences (liver failure; hepatocellular cancer conditions), and also for drug screening, diagnosis, target conditions of the function and gene mapping (for example of single-nucleotide polymorphisms). The chemical modification improves stability, activity,
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Best Local S
Matches 12
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11-MAR-2002;
26-MAR-2002;
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09-SEP-2002;
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05-AUG-2002;
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                                                                                                                                                                                                                                  Example 3; SEQ ID NO 1316; 183pp; English
                                                                                                                                                                                                                                                                           New double-stranded short modified pyrimidine bases
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2002WG-US009187.

; 2002US-0386782P.

; 2002US-0401104P.

; 2002US-0406784P.

; 2002US-0408378P.

; 2002US-0408379.

; 2003US-0440129P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cancer; cirrhosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RNA;
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                                                                                                                                                                                                                                                                           interfering nucleic useful for treating
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Pred. No.
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infection with hepatitis
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RESULT 25
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Best Local S
Matches 12
acids (siNA) that inhibits replication of hepatitis C virus (HCV), where one strand is an antisense strand (ASS) that is complementary to (part of) an HCV RNA (portion) and a sense strand (SS) that is complementary to (part of) and where most of the pyrimidine nucleotides comprise a sugar modification. The invention may allow development of compounds with virucide, antiinflammatory, hepatotropic or cytostatic activities by modulation (inhibition) of expression or activity of HCV RNA, by RNA interference. The siNA's of the invention may be used to inhibit replication of HCV, in cells, tissue explants or organisms, for treating HCV infection and its consequences (liver failure; hepatoccellular cancer and cirrhosis), and also for drug screening, diagnosis, target
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11-MAR-2002;
26-MAR-2002;
06-JUN-2002;
05-AUG-2002;
29-AUG-2002;
05-SEP-2002;
09-SEP-2002;
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conserved
strains.
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hepatotropic; cytos
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12; Conserv
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pyrimidine
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2002WG-US0091B7
2002US-0386782P
2002US-0401104P
2002US-0406784P
2002US-0406378P
2002US-0409293P
2003US-0440129P
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Best Local (
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This invention relates to novel double-stranded short interfering nucleic acids (siNA) that inhibits replication of hepatitis C virus (HCV), where one strand is an antisense strand (ASS) that is complementary to (part of) an HCV RNA (portion) and a sense strand (SS) that is complementary to ASS, and where most of the pyrinidine nucleotides comprise a sugar modification. The invention may allow development of compounds with virucide, antiinflammatory, hepatotropic or cytostatic activities by modulation (inhibition) of expression or activity of HCV RNA, by RNA interference. The siNA's of the invention may be used to inhibit replication of HCV, in cells, tissue explants or organisms, for treating HCV infection and its consequences (liver failure; hepatocellular cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-FEB-2002;
11-MAR-2002;
26-MAR-2002;
06-JUN-2002;
05-AUG-2002;
29-AUG-2002;
05-SEP-2002;
09-SEP-2002;
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                                                                                                                                                                                                                                                                   Example
                                                                                                                                                                                                                                                                                                             virus.
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hepatotropic; cytostatic;
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2002WO-US009187.
2002US-0386782P.
2002US-0401104P.
2002US-0406784P.
2002US-04008378P.
2002US-04008378P.
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interference;
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11-WAR-2002; 2002US-0363124P.
26-WAR-2002; 2002WG-03609187.
06-JUN-2002; 2002US-0386782P.
05-AUG-2002; 2002US-0401104P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
15-UAN-2003; 2003US-0440129P.
               This invention relates to novel double-stranded short interfering nucleic acids (siNA) that inhibits replication of hepatitis C virus (HCV), where one strand is an antisense strand (ASS) that is complementary to (part of) an HCV RNA (portion) and a sense strand (SS) that is complementary to ASS, and where most of the pyrimidine nucleotides comprise a sugar modification. The invention may allow development of compounds with virucide, antiinflammatory, hepatotropic or cytostatic activities by modulation (inhibition) of expression or activity of HCV RNA, by RNA modulation (inhibition)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ç;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 14;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0 T; 3 U;
invention may be used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 10; I
                                                                                                                                                                                                                                                                                                                                                                                                                                      Morrissey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0 Other;
                                                                                                                                                                                                                                                                                                             acid comprises sugar-
infection with hepatitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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ASS, and where most of the pyrimidine nucleotides comprise a su modification. The invention may allow development of compounds virucide, antiinflammatory, hepatotropic or cytostatic activiti

sugar

This invention relates to novel double-stranded short interfering nucleic acids (siNA) that inhibits replication of hepatitis C virus (HCV), where one strand is an antisense strand (ASS) that is complementary to of part of an HCV RNA (portion) and a sense strand (SS) that is complementary to

New double-stranded modified pyrimidine

short

interfering nucleic useful for treating

acid comprises sugar-infection with hepatitis

a

Example 3;

SEQ

ID NO 1288;

183pp; English

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Best Local S
Matches 12
                                                                                                                                                                                                                                                                                     20-FEB-2002;
11-MAR-2002;
26-MAR-2002;
06-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 replication of HCV, in cells, tissue explants or organisms, for treating HCV infection and its consequences (liver failure; hepatocellular cancer and cirrhosis), and also for drug screening, diagnosis, target identification and validation, genetic engineering, pharmacogenomics, studying gene function and gene mapping (for example of single-nucleotide polymorphisms). The chemical modification improves stability, activity, cellular uptake and/or binding affinity. The slNA can be directed to conserved regions of HCV genes, so are active against many different
                                                                                                                                                                                                                                                                                                                                                                                                                                        short interfering nucleic acid; siNA; virus hepatitis C virus; HCV; sugar modification; hepatotropic; cytostatic; RNA interference; hepatocellular cancer; cirrhosis; ss.
                                                                                                                                                           WPI; 2003-689778/65.
                                                                                                                                                                                                                                                     05-SEP-2002;
                                                                                                                                                                                                                                                                29-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19 BP; 3 A; 7 C; 5 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                15-JAN-2003;
                                                                                                                                                                                                                                                                                                                                              20-FEB-2003; 2003WO-US005043.
                                                                                                                                                                                                                                                                                                                                                                                            WO2003070750-A2
                                                                                                                                                                                                                                                                                                                                                                                                                  Hepatitis C virus.
                                                                                                                                                                                                         (SIRN-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hepatitis C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADF52698
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                139 ACGAGGTTGCGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12;
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                                                                                                                                                                                                         SIRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACGAGGUUGCGACC
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                                                                                                                                                                                                                            2002US-0363124P.
2002US-0386782P.
2002US-0386782P.
2002US-0401104P.
2002US-0406784P.
2002US-0408378P.
2002US-0408378P.
2002US-0408378P.
2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               virus siNA antisense strand SeqID1288
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                                                                                                                                                                                                                                                                                                                        2002US-0358580P
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                                                                                                                                                                                                         THERAPEUTICS INC
                                                                                                                                                                                 Beigelman L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4.4%;
85.7%;
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                                                                                                                                                                                 Macejak D,
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Pred.
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                                                                                                                                                                                 Morrissey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 10;
7.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 replication inhibition;
virucide; antiinflammatory;
HCV infection; liver failure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      <u>.</u>
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RESULT 29
ADF52007/c
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Best Local
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05-SEP-2002;
09-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                  26-MAR-2002;
06-JUN-2002;
05-AUG-2002;
 This invention relates to novel double-stranded short interfering nucleic acids (siNA) that inhibits replication of hepatitis C virus (HCV), where one strand is an antisense strand (ASS) that is complementary to (part of) an HCV RNA (portion) and a sense strand (SS) that is complementary to ASS, and where most of the pyrimidine nucleotides comprise a sugar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modulation (inhibition) of expression or activity of HCV RNA, by RNA interference. The siNA's of the invention may be used to inhibit replication of HCV, in cells, tissue explants or organisms, for treating HCV infection and its consequences (liver failure; hepatocellular cancer and cirrhosis), and also for drug screening, diagnosis, target identification and validation, genetic engineering, pharmacogenomics, studying gene function and gene mapping (for example of single-nucleotide polymorphisms). The chemical modification improves stability, activity, cellular uptake and/or binding affinity. The siNA can be directed to conserved regions of HCV genes, so are active against many different
                                                                                                                                                                New double-stranded modified pyrimidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hepatitis C virus short interfering
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADF52007
                                                                                                           Example 3;
                                                                                                                                                                                                                                                        Mcswiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-MAR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-FEB-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO2003070750-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            hepatocellular cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             hepatitis C virus; HCV; su hepatotropic; cytostatic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                              (SIRN-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 interfering nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             l Similarity
12; Conserv
                                                                                                                                                                                                                                                                                            SIRNA
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                                                                                                                                                                                                                                                        ŗ,
                                                                                                           SEQ
                                                                                                                                                                                                                                                                                                                          2002US-0358580P.
2002US-0363124P.
2002US-0386782P.
2002US-0386782P.
2002US-0401104P.
2002US-0406784P.
2002US-0408378P.
2002US-0408378P.
2002US-0408378P.
2003US-0440129P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2003WO-US005043.
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                                                                                                                                                                                                                                                        Beigelman
                                                                                                                                                                                                                                                                                            THERAPEUTICS INC
                                                                                                           ID NO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nucleic acid; siNA; virus HCV; sugar modification; static; RNA interference;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RNA; 19
                                                                                                                                                                l short
bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4.4%;
85.7%;
                                                                                                           597;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    7 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cirrhosis;
the pyrimidine nucleotides comprise a sugar
                                                                                                                                                                                                                                                      ŗ
                                                                                                         183pp,
                                                                                                                                                              interfering nucleic useful for treating
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                                                                                                                                                                                                                                                    Macejak
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Pred.
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                                                                                                                                                                                                                                                      Morrissey
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           virucide; antiinflammatory; HCV infection; liver failure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             replication
                                                                                                                                                              acid comprises infection with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             <u>,,</u>
                                                                                                                                                            sugar-
hepatitis C
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This invention relates to racids (siNA) that inhibits one strand is an antisense

on relates to novel double-stranded short interfering that inhibits replication of hepatitis C virus (HCV) s an antisense strand (ASS) that is complementary to

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NO 580;

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English.

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Best Local S
Matches 14
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11-MAR-2002;
26-MAR-2002;
06-JUN-2002;
05-AUG-2002;
29-AUG-2002;
29-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         virucide, antiinflammatory, hepatotropic or cytostatic activities by modulation (inhibition) of expression or activity of HCV RNA, by RNA interference. The siNA's of the invention may be used to inhibit replication of HCV, in cells, tissue explants or organisms, for treating HCV infection and its consequences (liver failure; hepatocellular cancer and cirrhosis), and also for drug screening, diagnosis, target identification and validation, genetic engineering, pharmacogenomics, studying gene function and gene mapping (for example of single-nucleotide polymorphisms). The chemical modification improves stability, activity, cellular uptake and/or binding affinity. The slNA can be directed to conserved regions of HCV genes, so are active against many different
                                                                             New double-stranded modified pyrimidine
                                                                                                                                       Mcswiggen
                                                                                                                                                                                                                                                                                                                                 28-AUG-2003.
                                                                                                                                                                                                                                                                                                                                                       WO2003070750-A2
                                                                                                                                                                                                                                                                                                                                                                               Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                     hepatotropic; cytostatic; RNA inte
hepatocellular cancer; cirrhosis;
                                                                                                                                                                                                                                                                                                                                                                                                                            short interfering nucleic acid; siNA; virus hepatitis C virus; HCV; sugar modification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19
                                                                                                                                                                                                                                                                                                        20-FEB-2003; 2003WO-US005043
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADF51990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modification.
                                                                                                                                                                (SIRN-)
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                                                                                                                                                                SIRNA THERAPEUTICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ACGAGGTTGCGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ACGAGGTTGCGACC 152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     standard;
                                                                                                                                       ŗ,
                                                                                                                                                                                     2002WO-US009187.
2002US-0386782P.
2002US-0401104P.
2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
2003US-0440129P.
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                                                                                                                                                                                                                                                                      2002US-0358580P.
2002US-0363124P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                        Beigelman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention may allow development of compounds with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A
                                                                                                                                                                                                                                                                                                                                                                                                                                                               short interfering nucleic acid
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                                                                              short
bases
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                                                                             interfering nucleic useful for treating
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                 interference;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7.5e+03;
hes 0;
                                                                                                                                        Morrissey
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                                                                                                                                                                                                                                                                                                                                                                                                                  HCV
                                                                                                                                                                                                                                                                                                                                                                                                                                      replication inhibition;
                                                                              acid comprises infection with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                 infection; liver failure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                sense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                               antintlammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                strand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
                                                                              sugar-
hepatitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                SeqID580.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
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CCX88X7777XRXX8888888888888X88X88XX8XX8XXAXX
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ADF52690
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cof) an HCV RNA (portion) and a sense strand (SS) that is complementary to RAS, and where most of the pyrimidine nucleotides comprise a sugar modification. The invention may allow development of compounds with Virucide, antinflammatory, hepatotropic or cytostatic activities by modulation (inhibition) of expression or activity of HCV RNA, by RNA interference. The siNA's of the invention may be used to inhibit replication of HCV, in cells, tissue explants or organisms, for treating HCV infection and its consequences (liver failure; hepatocellular cancer and cirrhosis), and also for drug screening, diagnosis, target identification and gene mapping (for example of single-nucleotide polymorphisms). The chemical modification improves stability, activity, cellular uptake and/or binding affinity. The siNA can be directed to creains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                       20-FEB-2002; 2002US-0358580P.
11-WAR-2002; 2002US-0363124P.
26-MAR-2002; 2002WS-0369187.
06-JUN-2002; 2002US-0386782P.
05-AUG-2002; 2002US-0401104P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378R-
09-SEP-2002; 2002US-0409293P.
15-JAN-2003; 2003US-0440129P.
                                                              New double-stranded modified pyrimidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                        short interfering nucleic acid; siNA; virus replication inhibition; hepatitis C virus; HCV; sugar modification; virucide; antiinflammatory; hepatotropic; cytostatic; RNA interference; HCV infection; liver failur
                                                                                                          WPI; 2003-689778/65.
                                                                                                                                                                                                                                                                                                                                   20-FEB-2003; 2003WO-US005043
                                                                                                                                                                                                                                                                                                                                                               28-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                     Hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                               hepatocellular cancer; cirrhosis; gs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hepatitis C virus siNA antisense strand SeqID1280.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADF52690
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19
                                                                                                                                                                 (SIRN-) SIRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14;
                                                                                                                              ı,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ACGAGGTTGCGACC 152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 4 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                 THERAPEUTICS INC
                                                                                                                                  Beigelman L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4.4%;
                                                               l short
bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              6 C; 6 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N
                                                             interfering nucleic useful for treating
                                                                                                                                  Macejak
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 14;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                  'n
                                                                                                                                  Morrissey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         7.5e+03;
                                                              acid comprises infection with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                              sugar-
hepatitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0,
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Example 3; SEQ

ij

1280;

183pp; English

invention

relates to ö

novel double-stranded short interfering nucleic

Example 3;

SEQ

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592; 183pp; English

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RESULT 32
ADF52002/c
ID ADF52
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cc acids (siNA) that inhibits replication of hepatitis C virus (HCV), where consistend is an antisense strand (ASS) that is complementary to (part of) an HCV RNA (portion) and a sense strand (SS) that is complementary to ASS, and where most of the pyrinidine nucleotides comprise a sugar completion. The invention may allow development of compounds with virucide, antiinflammatory, hepatotropic or cytostatic activities by modulation (inhibition) of expression or activity of HCV RNA, by RNA crimterference. The siNA's of the invention may be used to inhibit replication and its consequences (liver failure; hepatocellular cancer and cirrhosis), and also for drug screening, diagnosis, target (identification and validation, genetic engineering, pharmacogenomics, studying gene function and gene mapping (for example of single-nucleotide polymorphisms). The chemical modification improves stability, activity, cellular uptake and/or binding affinity. The siNA can be directed to conserved regions of HCV genes, so are active against many different
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                               11-MAR-2002;
26-MAR-2002;
06-JUN-2002;
                                     New double-stranded modified pyrimidine
                                                                                                                                                                           05-SEP-2002;
09-SEP-2002;
                                                                                                                                                                                                      05-AUG-2002;
29-AUG-2002;
                                                                               WPI; 2003-689778/65
                                                                                                        Mcswiggen J,
                                                                                                                                                                                                                                                                         20-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                       short interfering nucleic acid; siNA; virus hepatitis C virus; HCV; sugar modification; hepatotropic; cytostatic; RNA interference;
                                                                                                                                                                                                                                                                                                    20-FEB-2003; 2003WO-US005043
                                                                                                                                                                                                                                                                                                                                                                                  Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hepatitis C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADF52002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADF52002 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-FEB-2004
                                                                                                                                     (SIRN-)
                                                                                                                                                                                                                                                                                                                                                                                                             nepatocellular cancer; cirrhosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ò
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           139 ACGAGGTTGCGACC 152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12;
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                                                                                                                                    SIRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4.4%;
Similarity 85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ACGAGGUUGCGACC
                                                                                                                                                           2002US-0363124P.
2002US-0S09187.
2002US-0386782P.
2002US-0401104P.
2002US-0406784P.
2002US-0408378P.
2002US-0408378P.
2003US-0408378P.
2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             virus short interfering nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 3 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                         2002US-0358580P.
                                                                                                        Beigelman
                                                                                                                                    THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RNA; 19
                                      short
bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           7 C; 6 G; 0 T;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16
                                     interfering useful for t
                                                                                                        Macejak
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                        Ò
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3 U; 0 Other;
                                   g nucleic
treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 10;
                                                                                                      Morrissey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                         replication inhibition; virucide; antiinflammature HCV infection; liver fa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             acid sense strand SeqID592
                                      acid comprises infection with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                     sugar-
hepatitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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ADF51989/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention relates to novel double-stranded short interfering nucleic CC acids (slNA) that inhibits replication of hepatitis C virus (HCV), where CC one strand is an antisense strand (ASS) that is complementary to (part CC ASS, and where most of the pyrimiddine nucleotides comprise a sugar modification. The invention may allow development of compounds with CC virucide, antiinflammatory, hepatotropic or cytostatic activities by modulation (inhibition) of expression or activity of HCV RNA, by RNA CC interference. The slNA's of the invention may be used to inhibit CC replication of HCV, in cells, tissue explants or organisms, for treating CC HCV infection and its consequences (liver failure, hepatocellular cancer and cirrhosis), and also for drug screening, diagnosis, target cancer finction and validation, genetic engineering, pharmacogenomics, to studying gene function and gene mapping (for example of single-nucleotide CC polymorphisms). The chemical modification improves stability, activity, activity, collular uptake and/or binding affinity. The siNA can be directed to conserved regions of HCV genes, so are active against many different
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local
                                                                                                                                                                                                          11-MAR-2002;
26-MAR-2002;
06-JUN-2002;
                                                                                                                                                                  05-AUG-2002;
29-AUG-2002;
05-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                            short interfering nucleic acid; siNA; virus hepatitis C virus; HCV; sugar modification; hepatotropic; cytostatic; RNA interference;
          New double-stranded short interfering nucleic modified pyrimidine bases useful for treating
                                                                                                                                                                                                                                                       20-FEB-2002;
                                                                                                                                                                                                                                                                                   20-FEB-2003;
                                                                                                                                                                                                                                                                                                                                                                    Hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hepatitis C virus short interfering nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADF51989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                               28-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADF51989;
                                                                                                              (SIRN-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         139
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14;
                                                                                                             SIRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ACGAGGTTGCGACC 5
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                                                                                                                                     2002US-0358580P.
2002US-0363124P.
2002US-0386782P.
2002US-0386782P.
2002US-0401104P.
2002US-0406784P.
2002US-04003784P.
2002US-04003789.
2002US-04003789.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4.4%;
ilarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 3
                                                                                                                                                                                                                                                                                 2003WO-US005043.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                               Beigelman
                                                                                                            THERAPEUTICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Α.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            6 C; 7 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                cirrhosis;
                                                                               ۲
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                               Macejak
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 14;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                               á
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 10;
. 7.5e+03
                                                                               Morrissey
                                                                                                                                                                                                                                                                                                                                                                                                                            replication inhibition; virucide; antiinflammat
                                                                                                                                                                                                                                                                                                                                                                                                              HCV infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      acid sense strand
          acid comprises infection with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                           antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                              liver failure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
          sugar-
hepatitis C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SeqID579.
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RESULT 34
ADP05478/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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                                                                                                                                                                                                                Determining genotype or mutant of target nucleic acid by selecting genotype or mutant sequence, amplifying nucleic acid having specific sequence, hybridizing probe to nucleic acid, detecting nucleic acid, analyzing hybridization pattern.
                                                                                                                                                                                                                                                                                                                                WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                   27-SEP-2002; 2002JP-00284453
                                                                                                                                                                                                                                                                                                                                                                                                                                                              27-SEP-2002; 2002JP-00284453
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ss; probe; hepatitis C virus; infection; hepatitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hepatitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-APR-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     JP2004113195-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADP05478;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADP05478 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           conserved
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
                                                                                                                                                                                                                                                                                                                                2004-360138/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14 ACGAGGTTGCGACC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3/ SEQ ID NO 579; 183pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           regions of HCV genes, so are active against many different
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 4 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   probe #6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5 C; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4.48;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 10;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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The invention relates to a method of determining genotype or mutant of target nucleic acid comprising selecting the sequence characteristic of the genotype or mutant, amplifying the nucleic acid containing the characteristic sequence, allowing a probe specific to the genotype or mutant to hybridise the amplified nucleic acid, detecting the nucleic acid hybridised to the probe and analysing the hybridisation pattern.

for determining the

genotype or

mutant

of.

target nucleic

of. 얁

Example;

SEQ ID NO 6;

41pp; Japanese.

ព័ត្តព្រះពិត្តព្រះក្នុង ខេត្តព្រះក្នុង ខេត្តព្រះកិច្ចព្រះក្នុង ខេត្តព្រះក្រុង ខេត្តព្រះក្រសាធិបាន ខេត្តព្រះក្រុង ខេត្តព្រះក្រុង ខេត្តព្រះក្រុង ខេត្តព្រះក្រង ខេត្តព្រះក្រុង ខេត្តព្រះក្រង ខេត្តព្រះក្រុង ខេត្តព្រះក្រុង ខេត្តព្រះក្រុង ខេត្តព្រិត ខេត្ត

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RESULT 35
AAV09175/c
ID AAV091
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Best Local
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This is a novel oligomer comprising 20 covalently linked nucleotides which bind to the 5' Cap region of the interleukin-1 receptor (ILIR) mRNA. Expression of ILIR, in cells and tissues can be modulated by compositions comprising oligomers which are able to specifically hybridise with target areas of its encoding sequence. The composition cape used for treatment of disease in humans caused by excessive receptor expression, e.g. inflammation. When labelled they can be used diagnostically to determine overexpression of ILIR, also to determine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           acid, preferably hepatitis C virus. The method is useful for immobilising an nucleic acid fragment to a solid support. The method is useful for purifying and amplifying desired viral genome, preferably hepatitis B virus. The method is useful for amplifying desired nucleic acid and for discriminating the genotype and mutant of hepatitis B virus. The method is useful in detecting the genotype of patient infected with viral hepatitis, estimating the illness condition of patient, and predicting the treatment for the infection. The method efficiently and simply determines the genotype or mutant of target nucleic acid. The method is simple and rapid for discriminating genotype and mutant of hepatitis B virus at higher rate. The present sequence represents a hepatitis C virus
                                                                                                                                                                                                                                                                           2'-substituted oligonuclectide(s) specific for interleukin-1 retype I - used to modulate expression and detect overexpression
                                                                                                                                                                                                             Example 5;
                                                                                                                                                                                                                                                                                                                                                  WPI; 1998-018646/02
                                                                                                                                                                                                                                                                                                                                                                                            Miraglia L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-MAY-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-NOV-1997.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Type I interle
inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Phosphorothiate
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAV09175
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                                                                                                                                                                                                                                                                                                                                                                                                                                            (ISIS-) ISIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  interleukin-1
mation; 88; 5'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACGAGGTTGCGACC 152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ACGAGGTTGCGACC
                                                                                                                                                                                                             Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP; 3 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                            Bennett
                                                                                                                                                                                                                                                                                                                                                                                                                                            PHARM INC.
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/*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   97WO-US007147
                                                                                                                                                                                                           19; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   note= "Phosphorothioate"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.4%;
                                                                                                                                                                                                                                                                                                                                                                                            ÇF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               receptor; IL1R; human; IL1 protein; hybridisation; Cap region; phosphorothioate linkage.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        σı
                                                                                                                                                                                                                                                                                                                                                                                               Dean
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sequence
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                                                                                                                                                                                                                                                                                                                                                                                          Geiger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 12;
7.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   internucleotide linkage"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8054 targeting ILIR mRNA
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                           Best Loc
Matches
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Matches
                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                              AAX91991-X97517 represent PCR primers used to amplify open reading frames and other nucleic acid sequences from the genome of Chlamydia pneumoniae (see AAX91990). C. pneumoniae causes respiratory disease such as pneumonia and bronchitis and is thought to be a contributing factor in heart disease, sarcoidosis, sinusitis, purulent otitis media, erythema nodosum or pharyngitis. The polypeptides encoded by the open reading frames of the C. pneumoniae genome (see AAX34584- AAX35879) can be used in immunogenic compositions as vaccines. Vectors containing C. pneumoniae especially where the vector directs the expression of a neutralising epitope of C. pneumoniae
                                                                            Sequence
                                                                                                                                                                                                                                                                                Genome sequence of Chlamydia pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                    21-NOV-1997;
04-NOV-1998;
                                                                                                                                                                                                                                                       Page 1575; Disclosure; 1912pp; English.
                                                                                                                                                                                                                                                                                                                                   Griffais
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9927105-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sinusitis; purulent otitis media; ery neutralising epitope; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Respiratory disease; pneumonia; bronchitis; heart disease; sinusitis; purulent otitis media; erythema nodosum; pharvno
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 2 A; 7 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                          WPI; 1999-357842/30.
                                                                                                                                                                                                                                                                                                                                                                                                                        20-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chlamydophila pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primer used to amplify an ORF of Chlamydia pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAX93253;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAX93253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         localisation and distribution of this expression for research, or therapeutic purposes
                                                                                                                                                                                                                                                                                                                                                         (GEST ) GENSET
                                      Local Similarity
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  135
                           14;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                   R
 GGAGACGAGGTTGC 148
                                                                           20 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GCCGCCGCAGCCCT 1
                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                  98US-0107078P
                                                                                                                                                                                                                                                                                                                                                                                                                        98WO-IB001890
                                                                           4 A; 9 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 entry)
                                      100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
                                                    4.48;
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                                   Score 14;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14; DB; Pred. No. 7.5
                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                erythema nodosum; pharyngitis; vaccine;
                                     DB 2; Le
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 2;
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                                                 Length 20;
                         Indels
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RESULT 37
AAC67142/c
ID AAC671
XX AAC671
XX AAC671
XX AAC671
XX Human
DT 03-APR
XX Phomo 8
XX Phomo 8
XX US6165
XX TATE
PT Crans
PT Cr
 RESULT 38
ABT0851
ID ABT08
XX ABT08
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention provides antisense oligonucleotides with phosphorothicate backbones directed at the human E2F transcription factor 3 (E2F-3) coding sequences. These can be used in the therapy of diseases which can be treated by modulating E2F-3 expression and to prevent infection, inflammation and tumour formation
                                                           virucide; tranquilizer; anticonvulsant; osteopathic; analgesic; antiparkinsonian; dermatological; antiinfertility; cerebroprotective; antiaddictive; PCR; primer; probe; ss.
                                                                                                                                                 Human; NOVX; single nucleotide polymorphism; SNP; anti-HIV; cytostatic; antiarteriosclerotic; antidiabetic; antiasthmatic; antiinflammatory;
                                                                                                                                                                                                                                                      28-NOV-2002
                                                                                                                                                                                                                                                                                                                                ABT08551 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel antisense compounds targeted to E2F transcription factor 3 diagnosis, prophylaxis and treatment of diseases associated with transcription factor 3 such as infection, inflammation or tumor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-101698/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-FEB-2000; 2000US-00513729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAC67142
                                                                                                             haemostatic; hypotensive; neuroprotective; anorectic; nootropic; antidepressant; immunosuppressive; antibacterial; antiparasitic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-FEB-2000; 2000US-00513729
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                03-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ISIS-) ISIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-DEC-2000
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                       sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      E2F transcription factor 3 mRNA antisense sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E2F transcription factor 3; antisense; E2F-3; orothioate backbone; infection; inflammation; I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Col 41-42; 41pp; English
                                                                                                                                                                                                              protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                  (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Α,
                                                                                                                                                                                                                                                                                                                            DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA;
                                                                                                                                                                                                            coding sequence NOVX PCR primer
                                                                                                                                                                                                                                                  entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              7 C; 10 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4.4%;
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                                                                                                                                                                                                                                                                                                                                ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               infection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 14;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 4; Le
. 7.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                            SEQ ID
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                                                                                                                                                                                                            NO:
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E2F
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ADJ34405
ID ADJ3
XX
AC ADJ3
AC ADJ3
AC BDG
DE Huma
XX
KW Huma
XX
KW Huma
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KW endc
KW endc
KW endc
KW prim
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PN WO20
XX
                                                                                                                                                                                                                                                                                                                          RESULT 39
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                                                                                                               Human; ss;
endocrine
                   WO2004000997-A2
                                                                                                                                                                      Human secreted
                                                                                                                                                                                                            06-MAY-2004
                                                                                                                                                                                                                                                  ADJ34405;
                                                                                                                                                                                                                                                                                      ADJ34405 standard;
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CAGCACCAGTGGAA

254

Conservative

Indels

<u>,,</u>

Gaps

0

CAGCACCAGTGGAA

RTQ

PCR;

NOVX; secreted protein; cancer; der; CNS disorder; inflammatory of real time quantitative PCR.

diabetes; obesity; disorder; gene ther

quantitative

(first

entry)

DNA;

20

protein NOV5

RTQ

PCR

primer #2.

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Query Match
Best Local
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19-JAN-2001;
28-FEB-2001;
20-APR-2001;
26-JUL-2001;
09-AUG-2001;
                                                                                                                               The present invention provides the protein and coding sequences of several novel human proteins, designated NOVX. These can be used in the treatment of diseases such as cancers, Hodgkin's disease, Von Hippel-Lindau syndrome, Alzheimer's disease, stroke, tuberous sclerosis, hypercalcaemia, Parkinson's disease, Huntington's disease, cerebral palsy, epilepsy, Lesch-Nyhan syndrome, multiple sclerosis, ataxia telangiectasia, leukodystrophies, addiction, anxiety, depression, pain, obesity, Crohn's disease, osteoporosis, inflammatory bowel disease, infertility, atherosclerosis, hypertension, scleroderma, haemophilia, diabetes, pancreatitis, autoimmune disease, arthritis, immunodeficiencies, HIV, viral, bacterial or parasitic infections, or graft-versus-host disease. The present sequence is an oligonucleotide used to isolate a coding sequence of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Shimkets I
Shenoy S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New isolated NOVX polypeptides and polynucleotides, useful for preventing, diagnosing or treating NOVX-associated disorders e diabetes, Crohn's disease, atherosclerosis, cancer, Huntington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI;
                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 286; 363pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rothenberg M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-DEC-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Taupier RJ,
Tchernev V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CURA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2002-666903/71.
                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURAGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RA, Patturajan M. Vernet CAM, Casman SJ,
, Spytek KA, Gangolli E, Miller C, Boldog
RJ, Kekuda R, Smithson G, Zerhusen BD, Li
V, Si J, Edinger S, Stone D, Sciore P,
                                                                                              20
                                                                                            ₽₽;
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2001US-0285189P.
2001US-0308039P.
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100.0%;
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                           Score 14;
Pred. No.
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                           DB 6; Le 7.5e+03;
                                                                                            0 Other;
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BD, Liu X, C
ore P, Millet
                                              Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   disorders e.g.
Huntington's disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Malyankar U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Li ...
Colman
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Query Match Best Local S Matches 14

Similarity

100.0%; 4.48;

Score 14; Pred. No. Mismatches

DB 12; 7.5e+03; 0

Length

0,

Gaps

0

14;

Conservative

0

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The invention relates to a new isolated polypeptide (designated NOVX) CC comprising one of 141 fully defined sequences, their mature forms, a protein comprising one or more conservative substitutions or having at CC least 95% identity to one of the 141 proteins. Also included are a CC composition comprising NOVX (or a NOVX nucleic acid molecule (NA)), a kit comprising the composition of NOVX in one or more containers, an isolated CC nucleic acid molecule encoding a NOVX protein, producing NOVX (comprising CC culturing a cell under conditions that lead to expression of the polypeptide, where the cell comprises a vector comprising NOVX (A), comprising culturing a nagent for use in the treatment of a pathology that is related to aberrant expression or physiological interactions of NOVX, screening for a modulator of activity of or latency or predisposition to a cell comprising the vector, an antibody that immunospecifically binds to preventing a pathology associated with NOVX, treating a pathological cell comprising the vector, an antibody that immunospecifically binds to NOVX, determining the presence or amount of NOVX or the nucleic acid molecule in a sample, and determining the presence of or predisposition to a disease associated with altered levels of expression of MOVX or the nucleic acid compositions of the present invention are useful for the diagnosis and treatment of disorders associated with aberrant expression or activity of the NOVX polypeptide, such as cancer, diabetes, obesity, and endocrine, cell compositions of the present invention are useful for the diagnosis and treatment of disorders associated with aberrant expression or activity of the NOVX polypeptide, such as cancer, diabetes, obesity, and endocrine, cell cancer and predictive medicine. The present sequence is an RTQ (real time quantitative) PCR primer for an mRNA encoding a NOVX protein.
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Prayaga bo,
Rothenberg ME, bull
Rothernev V
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17-JUN-2002;
19-JUN-2002;
17-JUL-2002;
17-JUL-2002;
12-AUG-2002;
13-AUG-2002;
14-AUG-2002;
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07-JUN-2002;
10-JUN-2002;
11-JUN-2002;
11-JUN-2002;
12-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-MAR-2002;
04-JUN-2002;
05-JUN-2002;
06-JUN-2002;
06-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New isolated NOVX polypeptides useful for treating, preventing and diagnosing pathological conditions with NOVX-associated disorders, as cancer, obesity, diabetes and inflammatory or CNS diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example D; SEQ ID NO 294; 418pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2004-082483/08.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Anderson DW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      on DW, Boldog FL, Burgess CE, Casman SJ, Edinger SR;
A, Ellerman K, Gerlach VL, Gorman L, Guo X, Gusev VY,
Macdougall JR, Malyankar UM, Millet I, Ort T, Paddigan
ja SK, Patturajan M, Pena CEA, Peyman JA, Rieger DK;
iberg ME, Sciore P, Shenoy SG, Smithson G, Spytek KA, (
21 RJ, Tchernev VT, Vernet CAM, Voss EZ, Zerhusen BD, (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURAGEN CORP
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2002US-0386041P

2002US-0386974P

2002US-0386974P

2002US-03870540P

2002US-03870540P

2002US-0387659P

2002US-0387549P

2002US-0387549P

2002US-0389723P

2002US-0389723P

2002US-0389729P

2002US-0389729P

2002US-0389729P

2002US-0389729P

2002US-0389729P

2002US-0389729P

2002US-0389742P

2002US-0398742P

2002US-0398743P

2002US-0403486P

2002US-0403748P

2002US-0403748P

2002US-0403748P
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Zhong
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Д.
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26-JUL-2001;
09-AUG-2001;
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19-JAN-2001;
28-FEB-2001;
28-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             human; NOVX; immunogen; vaccine; cancer; diabetes; Alzheimer's disease; Parkinson's disease; Huntington's disease; asthma; allergy; emphysema; bronchitis; autoimmune disease; graft-versus-host disease; arthritis; scleroderma; systemic lupus erychematosus; bacterial infection; cystic fibrosis; coronary artery disease; stroke; hypertension; myocardial infarction; haemophilia; idiopathic thrombocytopenic purpura hyperlipidaemia; obesity; cirrhosis; inflammatory bowel disease; crohn's disease; ulcers; muscular dystrophy; myasthenia gravis; endometriosis; paoriasis; alopecia; uveitis; amyotrophic lateral sclerosis; osteoporosis; osteoarthritis; liver disease; epilepsy; multiple sclerosis; anxiety; pain; fertility; glomerulonephritis; polycystic kidney disease; ss; PCR; primer.
                         (KEKU/)
(SMIT/)
(ZERH/)
(LIUX/)
(LIUX/)
(COLM/)
(TCHE/)
(SIJJ/)
(SIJJ/)
(STON/)
                                                                                                                                                                                                                                                                           (TILL)
                                                                                                                                                                                                                                                                                                                                                    (MILL,
                                                                                                                                                                                                                                                                                                                                                                             (SHEN/)
(SPYT/)
(GANG/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27-MAR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US2004018970-A1
                                                                                                                                                                                                                                                                                                                            (BOLD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-JUL-2004 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                40
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                                                                                                                                                                                                                                                                                                                                                                                                                                           SHIMKETS R A.
PATTURAJAN M.
VERNET C A M.
CASMAN S J.
MALYANKAR U M.
                                                                                                                                                                                                                                                                                                                     SHENOY S G.
SPYTEK K A.
GANGOLLI E A.
MILLER C E.
BOLDOG F L.
EDINGER S R
STONE D J.
SCIORE P.
MILLET I.
                                                                                                                                                                          KEKUDA R.
SMITHSON G.
ZERHUSEN B D.
LIU X.
                                                                                                                      COLMAN S D.
                                                                                                                                                                                                                                                                    TAUPIER R J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         7 CAGCACCAGTGGAA 20
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2001US-0262959P.

2001US-0272408P.

2001US-0279344P.

2001US-0285189P.

2001US-030039P.

2001US-0311266P.

2001US-03028248.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2002US-00107782
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20
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RESULT 41
AAL41635/c
ID AAL416
XX AAL416
XX AAL416
XX AAL416
XX AAL416
XX Human
DE Human
XX Human
XX Human
XX Human
XX Human
XX Colore
XW metast
XX AX Humo
OS Homo 8
XX WO2001
XX WO2001
XX WO2001
XX WO2001
XX WO2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to an isolated NOVX polypeptide. A therapeutic, CC i.e. the NOVX polypeptide, nucleic acid and antibody, is useful for CC manufacturing a medicament for treating a syndrome associated with a CC human disease, e.g. a NOVX-associated disorder. The NOVX polypeptides can be used as immunogens or as vaccines. The NOVX polypeptide, nucleic acid CC or antibody is useful for diagnosing, treating or preventing a NOVX-associated disorder, e.g. cancer, diabetes, Alzhaimer's disease, arthritis, autoimmune disease, graft-versus-host disease, arthritis, arthritis, autoimmune disease, graft-versus-host disease, arthritis, CC scleroderma, systemic lugus erythematosus, bacterial infections, cystic ciproderma, systemic lugus erythematosus, bacterial infections, cystic ciproderma, systemic lugus erythematosus, bacterial infections, cystic ciprosis, coronary artery disease, stroke, hypertension, myocardial CC infarction, haemophilia, idiopathic thrombocytopenic purpura, CC disease, ulcers, muscular dystrophy, myasthenia gravis, endometriosis, CC osteoporosis, osteoarthritis, liver disease, spliepsy, multiple CC osteoporosis, osteoarthritis, liver disease, spliepsy, multiple CC sclerosis, anxiety, pain, fertility, glomerulonephritis, or polycystic cuseful for determining the presence of or preddisposition to a disease CC associated with altered levels of the NOVX polypeptide or the nucleic acid molecule, or for screening for molecules that inhibit or enhance CC probes, in chromosome mapping, tissue typing, preventive medicine, or CC probes, in chromosome mapping, tissue typing, preventive medicine, or CC prominer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 14
                                                                                                                     Human; colon cancer; cytostatic; drug design; adenomatous polyp; colorectal carcinoma; high metastatic potential colon tumour; metastatic colon cancer; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                     Human colon cancer
                                                                                                                                                                                                                                             19-APR-2002
                                                                                                                                                                                                                                                                                                                           AAL41635 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New NOVX polypeptides and nucleic acid molecules, useful for diagnosing, preventing or treating NOVX-associated disorders e.g. cancer, diabetes, Alzheimer's disease, stroke, arthritis, hypertension or myocardial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shinkets RA, Patturajan M, Vernet CAM, Casman SJ, Malyankar UM;
Shenoy SG, Spytek KA, Gangolli EA, Miller CE, Boldog FL, Li L;
Taupler RJ, Kekuda R, Smithson G, Zerhusen BD, Liu X, Colman SD;
Tchernev VT, Si J, Edinger SR, Stone DJ, Sciore P, Millet I;
                                          WO200196523-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; SEQ ID NO 159; 240pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2004-122080/12.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           infarction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ROTH/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  241 CAGCACCAGTGGAA 254
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14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                7
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                                                                                                                                                                                                                                                                                                                                                                                                                              CAGCACCAGTGGAA 20
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20-DEC-2001

The invention comprises isolated human gene sequences and PCR primer sequences which can be used to detect single nucleotide polymorphism (SNPs). The DNA sequences of the invention are useful for detecting a

le nucleotide polymorphisms are useful for detecting S

SNPs

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Claim 2; SEQ

ID NO 1778; 529pp; Japanese.

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RESULT 42
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to methods for detecting a cancerous colon cell involving detecting at least one differentially expressed gene such as those given in AALA1595-AALA1611. This is useful for diagnosing colon cancer, in rational drug and therapy design, and for identifying additional genes linked to the development or inhibition of development of colon cancer. Examples of colon cancer which can be detected include adenomatous polyp, colorectal carcinoma, high metastatic potential colon tumours and metastatic colon cancer. The present sequence is a PCR prime.
                                                                 Polynucleotide for detecting single nucleotide polymorphisms existing human gene, contains isolated human gene having specified sequence.
                                                                                                                                                      11-DEC-2001; 2001JP-00377637
                                                                                                                                                                             11-DEC-2001; 2001JP-00377637
                                                                                                                                                                                                                              JP2003174883-A
                                                                                                                                                                                                                                                                         human; gene sequence; single nucleotide polymorphism; disease diagnosis; ss; PCR; primer.
                                                                                                                                                                                                                                                                                                               Human gene PCR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Detecting a cancerous colon cell, useful for diagnosing colon cancer for rational drug and therapy design, comprises detecting at least or differentially expressed gene product.
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                                                                                                                                (KAGA-)
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Pred. No.
                                                                                                                                JIGYODAN
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hes 0;
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RESULT 43
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AAQ34273/c
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                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to a polynucleotide isolated from a human gene and is useful for detecting a single nucleotide polymorphism in a human gene or for diagnosing of disease. The invention enables the detection of a single nucleotide polymorphism in a human gene. The present sequence represents a primer of the invention.
            Upstream PCR primer TGLA137UP1.
                                               25-MAR-2003
02-FEB-1993
                                                                                                    AAQ34273;
                                                                                                                                   AAQ34273 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                 Sequence 24 BP; 8 A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        human; single nucleotide polymorphism; SNP; ss; primer.
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                                                                                                                                                                                                                                                           276
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                                                                                                                                                                                                                                                         TTGTTAAGAAGCAG 289
                                                                                                                                                                                                                          TTGTTAAGAAGCAG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 7364; 2627pp; Japanese.
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(first entry)
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7.6e+03;
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RESULT 45
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                                                                                                                                                                            EST; ss; probe; expressed sequence tag; microarray; gene expression; genetic variation; biallelic marker; polymorphism; human;
05-JUN-2003.
                                                     US2003104410-A1
                                                                                                                                                           cross-species
                                                                                                                                                                                                                                                                 Human microarray DNA oligonucleotide SEQ ID NO 100737.
                                                                                                                                                                                                                                                                                                                             14-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                     ACK00756;
                                                                                                                                                                                                                                                                                                                                                                                                                                   ACK00756 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Polymorphic bovine DNA markers - used in mapping, and selective breeding.
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RESULT 46
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   or family members of a gene and a cross-species comparison. Each of the nucleic acids further comprises a tag sequence. The array of nucleic acid probes is useful in in situ hybridisation, in Southern, Northern or dotblot hybridisation to identify or detect the sequence or specific mutations of any gene, in mapping the 5' termini of many molecules by primer extensions or in screening cDNA or genomic libraries or subclones for additional subclones containing segments of DNA that have been isolated and previously sequenced. The sequence presented is one of the nucleic acid probes incorporated in the microarray. Note: The sequence data for this patent can also be obtained in electronic format directly from USPTO at sequence/sequence.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention discloses a microarray comprising a plurality of nucleic acid probes including one of 2,018,500 fully defined sequences, or its perfect match, perfect mismatch, antisense match or antisense mismatch. Also disclosed is a method of gene expression analysis. The array is used in monitoring gene expression levels by hybridisation to a DNA library, in analysis of genetic variation or in hybridisation of tag-labelled compounds. The nucleic acid probes are specifically designed for analysis of at least one target sequence. The method of analysis comprises hybridising at least one or more nucleic acids to at least two or more nucleic acid probes are attached to a solid support. The analysis comprises monitoring gene expression levels, identifying biallelic markers or polymorphisms,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New array of nucleic acid probes, useful for Southern, Northern or dot-blot hybridization sequence or specific mutations of any gene.
                                                                                                                                               EST; ss; probe; expressed seq
genetic variation; biallelic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 25 BP; 9
                                             05-JUN-2003.
                                                                                                                                                                                          Human microarray DNA oligonucleotide SEQ ID NO 86601
                                                                                                                                                                                                                          14-OCT-2003
                                                                                                                                                                                                                                                                                  ACI86610 standard; DNA; 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-567953/53
               15-MAR-2002; 2002US-00098263
                                                                           US2003104410-A1
                                                                                                      Homo sapiens
                                                                                                                                   стовв-вресіев
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (AFFY-) AFFYMETRIX INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-MAR-2001; 2001US-0276759P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-MAR-2002;
                                                                                                                                                                                                                                                                                                                                                                                       172
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                                                                                                                                                                                                                                                                                                                                                                                                                    14;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                               ACTGGTCTCTGGAG 185
                                                                                                                                                                                                                                                                                                                                                         ACTGGTCTCTGGAG 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO 100737; 9pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2002US-00098263.
                                                                                                                                                                                                                      (first entry)
                                                                                                                                   comparison.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7 C; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                               4.48;
                                                                                                                                            sequence tag; microarray; gene expression;
lic marker; polymorphism; human;
                                                                                                                                                                                                                                                                                                                                                                                                                  0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                Score 14;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                              DB 9; L.,
0. 7.6e+03;
0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 25
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                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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CC in monitoring gene expression levels by hybridisation to a DNA library.

CC in analysis of genetic variation or in hybridisation to a DNA library.

CC compounds. The nucleic acid probes are specifically designed for analysis compounds the ast one target sequence. The method of analysis comprises comprises are attached to a solid support. The analysis comprises mucleic acid probes and detecting the hybridisation. The nucleic acid crobes are attached to a solid support. The analysis comprises monitoring compression levels, identifying biallelic markers or polymorphisms, cor ramily members of a gene and a cross-species comparises monitoring cor family members of a gene and a cross-species comparises. Each of the nucleic acid crobes is useful in in situ hybridisation, in Southern, Northern or dotolot hybridisation to identify or detect the sequence or specific cutations of any gene, in mapping the 5' termini of mRNA molecules by primer extensions or in screening cDNA or genomic libraries or subclones containing segments of DNA that have been contained and previously sequenced. The sequence presented is one of the nucleic acid probes incorporated in the microarray. Note: The sequence contained in the microarray note format directly from USPTO at sequence can also be obtained in electronic format directly contained in electronic format directly contained in the microarray.
Sequence 25 BP; 7 A; 9 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention discloses a microarray comprising a plurality of nucleic acid probes including one of 2,018,500 fully defined sequences, or its perfect match, perfect mismatch, antisense match or antisense mismatch. Also disclosed is a method of gene expression analysis. The array is used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; SEQ ID NO 86601;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New array of nucleic acid probes, useful for Southern, Northern or dot-blot hybridization sequence or specific mutations of any gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-MAR-2001; 2001US-0276759P
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Matches
                                         Query Match
Best Local :
                                        Local
               172 ACTGGTCTCTGGAG 185
16
                                        Similarity
ACTGGTCTCTGGAG 3
                                 Conservative
                                       4.4%;
                                 <u>.</u>
                                        Score 14;
Pred. No.
                                 Mismatches
                                       DB 9; L
                                 0
                                 Indels
                                0;
                                Gaps
                                 0
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Length 25

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RESULT 47
ACK00130/c
Human microarray DNA oligonucleotide SEQ ID NO 100111.
                                         14-OCT-2003
                                                                                                                      ACK00130 standard; DNA; 25
                                       (first entry)
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16-MAR-2001; 2001US-0276759P 15-MAR-2002; 2002US-00098263 05-JUN-2003

Homo sapiens

US2003104410-A1

cross-species comparison.

EST; ss; probe; expressed sequence tag; microarray;
genetic variation; biallelic marker; polymorphism;

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RESULT 48
ADP15727
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CC perfect match, perfect mismatch, antisense match or antisense mismatch.

CC Also disclosed is a method of gene expression analysis. The array is used

CC in analysis of genetic variation or in hybridisation to a DNA library,

CC in analysis of genetic variation or in hybridisation of taglabelled

CC compounds. The nucleic acid probes are specifically designed for analysis

CC of at least one target sequence. The method of analysis comprises

CC hybridising at least one or more nucleic acids to at least two or more

CC nucleic acid probes and detecting the hybridisation. The nucleic acid

CC probes are attached to a solid support. The analysis comprises monitoring

CC or family members of a gene and a cross-species comparison. Each of the

CC nucleic acids further comprises a tag sequence. The array of nucleic acid

CC probes is useful in in situ hybridisation, in Southern, Northern or dot-

CC mutations of any gene, in mapping the 5' termini of mRNA molecules by

CC primer extensions or in screening cDNA or genomic libraries or subclones

CC for additional subclones containing segments of DNA that have been

CC isolated and previously sequenced. The sequence presented is one of the

CC nucleic acid probes incorporated in the microarray. Note: The sequence

CC data for this patent can also be obtained in electronic format directly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
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                             21-NOV-2002;
03-APR-2003;
                                                                                                                   10-JUN-2004
                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                               ss; diagnosis; non-blood disease; solid tumor; gene expression; peripheral blood mononuclear cell; renal cell carcinoma; prostate cancer; head/neck cancer; differential expression; probe.
                                                                                                                                                                                                                                                                              Renal cell carcinoma differentially expressed gene probe #2132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                     ADP15727 standard; DNA; 25
                                                                                                                                                  WO2004048933-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (AFFY-) AFFYMETRIX INC
(AMHP ) WYETH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      USPTO at segdata.uspto.goc/sequence.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        172
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO 100111; 9pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ACTGGTCTCTGGAG 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ACTGGTCTCTGGAG 185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                               2002US-0427982P.
2003US-0459782P.
                                                                                  2003WO-US037481
                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             discloses a microarray comprising a plurality of a nocluding one of 2,018,500 fully defined sequences,
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G
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 14;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 9; Lo. 7.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0 Other;
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            20-SEP-2002; 2002WO-US029971
                                                27-MAR-2003
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RESULT 49
ABQ80298/c
ID ABQ802
XX ABQ802
XX ABQ802
XX Primer
XX BSP; C 20-SEP
XX 20-SEP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                             Primer; PCR; amplify; regulation; cell growth; myocardial infarction; osteoblast differentiation; CD49c; CD90; telomerase; p21; p53; CBFA1; BSP; core binding factor 1; bone sialoprotein; degeneration; trauma; acute injury; cardiac; neurological; spinal cord injury; amyotrophic lateral sclerosis; Parkinson's disease; stroke; traumatic brain injury; Fabry Disease condition; brain tumour; metachromatic dystrophy; adrenal leukodystrophy; Canavan disease; pelizaeus Merzbacher; Nieman-pick; congestive heart failure; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relate to a method of diagnosing (M1) non-blood disease such as solid tumor by providing peripheral blood sample of human having non-blood disease, and comparing an expression profile of specific genes in the peripheral blood sample to reference expression profile of the genes, where each of the genes is differentially expressed in peripheral blood mononuclear cells (PBMCs) of patients having the disease as compared to PBMCs of normal humans. The method is useful for diagnosing non-blood disease such as solid tumor. The solid tumor is chosen from renal cell carcinoma (RCC), prostate cancer and head/neck cancer. The peripheral blood sample comprises enriched PBMCs. The peripheral blood sample is a whole blood sample (claimed). (M1) is useful for identifying genes that are differentially expressed in peripheral blood samples isolated at different stages of progression, development or treatment of RCC and/or other solid tumors. This sequence corresponds to a probe to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Diagnosing non-blood disease such as solid tumor, involves differential expression profile of specific genes in periph sample of subject with reference expression profile of spec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (BURC/)
(TREP/)
(DORN/)
                                                                                                                                                                                                                                                                                                                                         Primer Telo-1F.
                                                                                                                                                                                                                                                                                                                                                                            27-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABQ80298 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 25 BP; 2 A; 6 C; 10 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   detect a gene that is differentially expressed and of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; SEQ ID NO 2463; 350pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-460799/43.
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                                                                                                                               Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  58 GGGACGGCTCCTGC 71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14;
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BURCZYNSKI M I
TREPICCHIO W I
DORNER A.
STOVER J A.
SLONI D K.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                          (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7.6e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    detected by the method
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n peripheral blood
of specific genes.
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ADKODO ADKODO
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              telomerase. The expression of transcripts for telomerase, p21, p53, CBFA1 and BSP were determined using quantitative PCR. Cell populations which co-express CD49c and CD90 express telomerase at the level of approximately 13 transcripts/10 power 6 transcripts of 18S rRNA, and these populations continue to proliferate at a constant rate. Compositions containing the cell populations of the invention are useful in treating humans suffering from a degenerative, traumatic, acute injury, cardiac or neurological condition, e.g., such as spinal cord injury, amyotrophic lateral sclerosis, Parkinson's disease, stroke, traumatic brain injury, Fabry Disease condition, metachromatic dystrophy, adrenal leukodystrophy, Canavan disease, Pelizaeus Merzbacher, Nieman-pick or brain tumour, myocardial infarction or congestive heart failure. This primer is
                                                                                                                                                                                                                                                                                                                                                                                           cardiac-related transcription factor; myocardial infarction; congestive heart; quantitative PCR; primer; ss; human; 18s RNA; cell growth regulator; osteoblast differentiation regulator; te
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              determine the expression of transcripts encoding regulators of cell growth and osteoblast differentiation. These primers were used to test the cell populations of the invention which co-express CD49c, CD90 and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        myocardial infarction or congestive deposited under Genbank ID AF015950
                                                                                                                                                                                                                                                                                       US2004058412-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cardiant; homogenous cell population;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADK00061
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example
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                                HO TW,
                                                                                                                                                                                                                                                                                                                                        Homo sapiens
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                                                                                                                              20-SEP-2002; 2002US-00251685
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Matches 14
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                                                                            The invention describes a complex of protein-protein interactions Helicobacter pylori selected from 421 complexes given in the specification. The complex of protein-protein interactions are use screening for agents which modulate the interaction of proteins. Modulating compounds which binds to a targeted bacterial protein modulating compounds which binds to a targeted bacterial protein modulating compounds which binds to a targeted bacterial protein modulating compounds which binds to a targeted bacterial protein modulating compounds which binds to a targeted bacterial protein modulating compounds which binds to a targeted bacterial protein modulating compounds which binds to a targeted bacterial protein modulating compounds which binds to a targeted bacterial protein modulating compounds which binds to a targeted bacterial protein modulating compounds which binds to a targeted bacterial protein modulating compounds which binds to a targeted bacterial protein modulating compounds which binds to a targeted bacterial protein modulating compounds which binds to a targeted bacterial protein modulating compounds which binds to a targeted bacterial protein modulating compounds which binds to a targeted bacterial protein modulating compounds which binds to a targeted bacterial protein modulating compounds which binds to a targeted bacterial protein modulating compounds which believes the comp
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factor, useful for preparing a pharmaceutical
myocardial infarction in humans.
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                                                                                                                                                                                                                                                                                New complexes of protein-protein useful for identifying modulating
                                                                                                                                                                                                                                                                                                                                                                                                 Legrain
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 5;
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                                                                                                                                                                                                                    Page 510;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 8 A;
                                                                                                                                                                                                                                                                                                                                                                                                 Rain J,
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used for treating or preventing ulcers in a human or animal. This sequence represents a primer used to isolate polynucleotides encoding Helicobacter pylori proteins for studies on protein-protein interacti

interactions

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RESULT 52
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a novel expression vector comprising an nEGFP reporter gene linked downstream of a HIV long terminal repeat (LTR) where a Rev protein nucleolus transfer signal is linked downstream of the reporter gene. The vector of the invention may be useful for infecting a cell with HIV-1 and measuring the infectivity of HIV-1 in a cell, as well as for producing HIV-1 clones, which in turn may be useful for anti-HIV strategies such as vaccine development. The current sequence is that of the HIV-1 Rev NLS (-) oligonucleotide of the invention which is related to HIV-1 Rev protein nucleolus transfer signal peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel expression vector comprising nEGFP reporter gene linked downstream of HIV-long terminal repeat, and a Rev protein nucleolus transfer signal is linked downstream of reporter gene.
                                      ABC96919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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                                                                                           ABC96919 standard; DNA; 13
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                                                                                                                                                                                                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
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         This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 ABC99889, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073
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                                                                                                                                                                                                                                                                                                                                                                                                        Oligonucleotide SEQ ID NO 96935 for detecting
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                                                                                                                                                                                                                           (EPIG-)
                                                                                                                                                                                                                                                                                                                  WO200177384-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TAGAATTGTAGTG
                                                                                                                                                                                                    Piepenbrock C,
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 oligomers described in the
                                                                                                              96935;
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Pred. No.
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invention. NOTE:
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The present invention describes nucleic acids (A) that interact stably CC with a target sequence and contain at least one phosphoro(di)thioate CC link, having endonuclease activity. (A), and more generally any catalytic CC nucleic acid (A') that modulates expression of the oestrogen receptor CC in vivo or by transforming cells ex vivo and implanting treated cells, or CC for other conditions associated with levels of oestrogen receptor. CC Because of the high selectivity for targeted RNA, (A) can also be used to CC correlate inhibition of gene expression with alterations in phenotype, CC particularly for identification of therapeutic targets, and as research CC reagents (for RNA, in the same way that restriction endonucleases are CC used with DNA). The combination of modifications in (A) improves CC resistence to nucleases, binding affinity and/or activity. AAA23503 to AAA24747 represent oestrogen receptor hammerhead ribozyme sequences, and AAA2593 to AAA25992 represent their corresponding target sequences. CC AAA2474 b to AAA25992 represent toestrogen receptor hairpin ribozyme sequences. AAA2610 to AAA26218 represent other ribozyme sequences and CC Sequences. AAA2619 to AAA26211 represent other ribozyme sequences and CC antisense oligonucleotides used in the exemplification of the present
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AAA26133
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Best Local Similarity
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Reynolds M, Zwick
Matulic-Adamic J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 79; Page 99; 148pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-APR-1998;
23-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oestrogen receptor; c-raf; k-ras; bcl-2; ribozyme; cleavage; hammerhead ribozyme; hairpin ribozyme; antisense oligonucleotide; gene expression modification; cancer; phosphorothioate; endonuclease; anticancer; breast cancer; endometrium cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oestrogen
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Zwick M,
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98US-00103636
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ris T, Woolf T,
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Pred. No.
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Haeberli
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RESULT 59 AAZ62579/c

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Best Local Similarity
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                                                                                                                              The present invention relates to a method for analysing genetic predisposition to an autoimmune disease. The method comprises producing at least one amplification product from a polymorphic region associated with the disease, and testing it for hybridisation to probes specific fo susceptibility, resistance and neutrality. The method is used to determine susceptibility to insulin-dependent diabetes, but can also be applied to rheumatoid arthritis, anhylosing spondylitis, lupus and scleroderma. The present sequence is a probe for HIA (human leukocyte antigen) DQB1 gene detection. The HIA-DQB1 gene is known to be associate with insulin-dependent diabetes.
                                                                                               Sequence
                                                                                                                                                                                                                                                                                  Assessing genetic predisposition to autoimmune diseases, specifically type I diabetes, by detecting polymorphisms associated with susceptibility, resistance or neutrality.
                                                                                                                                                                                                                                                          Claim 10; Page 43; 57pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; HLA-DQB1; probe; autoimmune disease; ankylosing spondylitis; insulin-dependent diabetes; rheumatoid arthritis; lupus; scleroderma;
                                                                                                                                                                                                                                                                                                                                    WPI; 2002-454864/48.
                                                                                                                                                                                                                                                                                                                                                                                                           17-NOV-2000; 2000FR-00014896
                                                                                                                                                                                                                                                                                                                                                                                                                                    16-NOV-2001; 2001WO-FR003599
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CGCACGTCCTCCT 1
               CGCACGTCCTCCT 108
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Pred. No. 2.5e+
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RESULT 60
ABX00430/c
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Best Local
                                                                                                                                                                                                                                                                                                                           The present sequence represents the preferred target sequence of an enzymatic nucleic acid, especially a hammerhead ribozyme, which cleaves the Hepatitis C virus (HCV) RNA sequence at the base position given in the descriptor line. The HCV sequence was screened for optimal ribozyme target sites using a computer folding algorithm and regions of the mRNA which did not form secondary folding structures and contained potential ribozyme cleavage sites were identified. Ribozymes were synthesised to target these sites and their activities optimised by either varying the length of the binding arms or by modification to prevent degradation by nucleases. The ribozymes of the invention inhibit gene expression and/or viral replication, and are used to treat diseases associated with Hepatitis C virus (HCV) infection, e.g. cirrhosis, liver failure and hepatocellular carcinoma. The ribozymes may be used in combination with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel ribozymes for the treatment hepatitis C infection.
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18-SEP-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Enzymatic nucleic acid; hammerhead ribozyme; virus replication; cirrhosis; liver failure; hepatocellular carcinoma; interferon;
                                                                                                                                                                                                                                                                         Sequence 15 BP; 1 A; 9 C; 3 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1;
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23-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Substrate for HH ribozyme HCV-3349 which cleaves HCV RNA at nt.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-MAR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ62579 standard; RNA; 15 BP.
            Hepatitis C
                                         23-DEC-2002
                                                                   ABX00430;
                                                                                              ABX00430
                                                                                                                                                                                                                                                                                                     diseases,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (RIBO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-APR-1999;
                                                                                                                                                                                                                                                                                                                 nterferon to treat HCV infection,
                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                           129
                                                                                                                                                                15
                                                                                                                                                                                                                     13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Page 56; 123pp; English.
                                                                                              standard;
                                                                                                                                                                  TCGGGCGGAGACG 3
                                                                                                                                                                                          TCGGGCGGAGACG 141
                                                                                                                                                                                                                                                                                                     and cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mcswiggen JA,
              virus substrate #212 for HCV
                                                                                                                                                                                                                     Conservative
                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98US-0083217P.
98US-0100842P.
99US-00257608.
99US-00274553.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99WO-US009027.
                                                                                              RNA; 15
                                                                                                                                                                                                                                 4.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Roberts
                                                                                                                                                                                                                     0,
                                                                                                                                                                                                                                 Score 13;
Pred. No.
                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     of diseases
                                                                                                                                                                                                                                                                                                                    other infectious
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pavco
                                                                                                                                                                                                                                 DB 3; Le
            hammerhead ribozyme #212.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PA,
                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     and conditions related
                                                                                                                                                                                                                                              Length 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Macejak
                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                  diseases,
                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                  autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cleavage; cancer;
                                                                                                                                                                                                                     Gaps
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SXSXSSXXS
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Best Local S
Matches 13
                                                                                                                                                                                                                                                                                                                                                              (HP) motif where the binding arms comprise sequences complementary to one of the substrate sequences defined in the specification. The HCV ribozymes are useful for modulating the expression and/or replication of HCV. They can be used to treat cirrhosis, liver failure and/or hepatocellular carcinoma. The HCV infection in conjunction with one or more other drug therapies, particularly type I interferon, especially interferon alpha, beta or gamma or consensus interferon. The present sequence represents a substrate for a HCV hammerhead (HH) ribozyme. Note: some of the sequence data for this patent did not form part of the printed specification. The complete sequence data for this patent was obtained in electronic format directly from the USPTO web site at
                                                      25-MAR-2003
08-JAN-1993
                                                                                                   AAQ26476;
                                                                                                                                                                                                                                                                                                                          Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   enzymatic nucleic acid or ribozyme is in a hammerhead (HH) or hairpin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention specifically cleave
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 27; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cirrhosis, liver failure or hepatocellular carcinoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New ribozymes targeting RNA derived from hepatitis C virus inhibit viral replication and are useful to treat hepatitis C virus infections and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Blatt L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (MCSW/)
(ROBE/)
(PAVC/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US2002082225-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Enzymatic nucleic acid; RNA cleavage; Hepatitis C virus infection; HCV ribozyme; HCV expression; HCV replication; cirrhosis; virucide; liver failure; hepatocellular carcinoma; HCV infection; drug therapy; type I interferon; interferon alpha; interferon beta; cytostatic;
                                                                                                                               AAQ26476 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                       seqdata.uspto.gov/psipsDIDEntry.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      interferon gamma; consensus interferon; hepatotropic; antiinflammatory; substrate; hammerhead ribozyme; HH ribozyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MACE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (BLAT/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2002-617759/66.
                                                                                                                                                                                                                                     129
                                                                                                                                                                                                       15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BLATT L.
MCSWIGGEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MACEJACK D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ROBERTS B. PAVCO P A.
                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                   TCGGGCGGAGACG 141
                                                                                                                                                                                                      TCGGGCGGAGACG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mcswiggen
                                                                                                                                                                                                                                                                                                                        BP; 1 A; 9 C; 3 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention
                                                                                                                                                                                                                                                                 Conservative
                                                      (revised)
(first en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99US-00274553
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                                                      entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 on relates to enzymatic nucleic acids which RNA derived from Hepatitis C virus (HCV)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  JA,
                                                                                                                                                                                                                                                                                100.0%;
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                                                                                                                                В₽
                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                Score 13; Pred. No.
                                                                                                                                                                                                                                                                  Mismatches
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                                                                                                                                                                                                                                                                                2.5e+04;
                                                                                                                                                                                                                                                                                             DB 6;
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                                                                                                                                                                                                                                                                  <u>,</u>
                                                                                                                                                                                                                                                                                           Length 15;
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                                                                                                                                                                                                                                                                 Gaps
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PCR; polymerase chain reaction; amplify; class

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HLA DQB1; probe;

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Probe DB69

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RESULT 62
AAQ26453
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The sequences given in AAQ26461-81 are probes which were used within the scope of the invention to type class II HLA DQB1 alleles. These probes were used to screen sequences amplified from the DQB1 gene second exon sequence. This method could be use to identify new DQB1 alleles. This method provides a simple, rapid and precise system for DQB1 typing, including those alleles which cannot be distinguished by serological methods. The presence or absence of a particular HLA DQB1 allele serves as an indicator of susceptibility to insulin-dependent diabetes mellitus organ transplantation and in the study of the molecular basis of disease susceptibility. Moreover, samples from unusual sources, eg ancient DNA or forensic samples, can be typed, even when the DNA sample is degraded or only present in an small amount. (Updated on 25-MAR-2003 to correct PNA field)
  21-DEC-1990;
                          20-DEC-1991;
                                                     09-JUL-1992.
                                                                              WO9211389-A1
                                                                                                        Synthetic.
                                                                                                                              PCR; polymerase chain reaction; amplify; class II HLA DQB1; probe; insulin-dependent diabetes mellitus; IDDM; forensics; ss.
                                                                                                                                                                        Probe DB69
                                                                                                                                                                                                 25-MAR-2003
08-JAN-1993
                                                                                                                                                                                                                                      AAQ26453;
                                                                                                                                                                                                                                                                AAQ26453 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 30; 37pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Erlich HA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1992-250108/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-DEC-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9211389-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  insulin-dependent diabetes mellitus; IDDM; non-coding strand; forensics;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (HOFF ) HOFFMANN LA ROCHE & CO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ldentity, and
                                                                                                                                                                                                                                                                                                                                                          96
                                                                                                                                                                                                                                                                                                                                                                                  13;
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                                                                                                                                                                                                                                                                                                                               CGCACGTCCTCCT 16
                                                                                                                                                                                                                                                                                                                                                          CGCACGTCCTCCT 108
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                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 3 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    for typing HLA DQB1 alleles -
d for studying disease suscepti
                                                                                                                                                                                                (revised)
(first entry)
90US-00632180.
                          91WO-US009796.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          90US-00632180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   91WO-US009796
                                                                                                                                                                                                                                                                                                                                                                                               4.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                    7 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A DQB1 alleles - for tissue typing, determining disease susceptibility.
                                                                                                                                                                                                                                                                                                                                                                                  0,
                                                                                                                                                                                                                                                                                                                                                                                               Score 13;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                          DB 2;
                                                                                                                                                                                                                                                                                                                                                                                               2.5e+04;
                                                                                                                                                                                                                                                                                                                                                                                                          Length 16;
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PACE SERVING SOLVE SERVING SER

Claim 10; Page 220; 345pp; Japanese.

transplanting

between them

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RESULT 63
ABL31182/c
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Best Local
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Human leukocyte antigen (HLA) tindividuals e.g. by determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; human leukocyte antigen; HLA; genotype; polymorphism; immunogenetic; transplantation; genetic disease; ss.
                                                                                                                                                                                                                                                                             01-JUN-2000; 2000JP-00164798
                                                                                                                                                                                                                                                                                                                               01-JUN-2001; 2001WO-JP004662
                                                                                                                                                                                                                                                                                                                                                                                        06-DEC-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human
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                                                                                    WPI; 2002-122074/16.
                                                                                                                                          Inoko H,
                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200192572-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABL31182;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Erlich HA,
                                                                                                                                                                                             (NISN ) NISSHINBO IND INC. (SYST-) SYSTEM RES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           field.)
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                                                                                                                                     Kagiya T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bugawan
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4.1%;
                                                                                                                                          Ichihara
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16
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Pred. No.
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  ping, useful for judging HLA genotypes immunogenetic differences when
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                                                                                                                                          Matsumura
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                                                                                                                                     Nishida
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OCCONTRACTION OF THE RESERVE TO A SECTION OF THE SE
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ABV78964
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                                                                                                                                                                                Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; human testis expressed Patched like protein; testis; adrenal; liv. male germ cell development; bone marrow; brain; kidney; lung; plap prostate; skeletal muscle; colon; male infertility; cancer; ss.
  The present invention relates to human testis expressed Patched 1 protein (HTPL, see ABV78759 to ABU78752 and ABB98519 to ABB98520) has two isoforms, with a few single base pair differences between two. One of the single base pair changes introduces a premature s
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o isoforms, with a
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shares an overall structure organisation with the Patched protein. The
construction of the patched protein and is a potential tumour suppressor. HTPL is
consider that of Patched, and is a potential tumour suppressor. HTPL is
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23-MAY-2001;
09-OCT-2001;
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                                                                                                   Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL.
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2001US-00327898P.
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protein

The present

invention relates to human testis expressed PL, see ABV78759 to ABV78762 and ABB98519 to

Patched like ABB98520). HTPL

718pp;

English

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RESULT 66
ABV78968
ID ABV78968
XX ABV78968
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30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
23-MAY-2001; 2001WO-US000669.
23-MAY-2001; 2001WS-03864761.
09-OCT-2001; 2001US-0327898P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    male germ prostate;
                                                                             Novel isolated human testis expressed Patched like protein (HTPL), us for identifying agonist and antagonist and specific binding partners, for treating subjects having defects in HTPL.
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Pred. No.
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2; Page 91;

718pp; English

useful rs, and

2002-676582/73.

Novel isolated human testis expressed Patched like protein (HTPL), use for identifying agonist and antagonist and specific binding partners, for treating subjects having defects in HTPL.

useful rs, and

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RESULT 67
ABV78967
ID ABV78
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 13; Conserv
                                                                                                                                                                 30-JAN-2001;
23-MAY-2001;
09-OCT-2001;
                                                                                                                                                                                                                                           30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                            Human; gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; human testis expressed Patched like protein; testis; adrenal; liver; male germ cell development; bone marrow; brain; kidney; lung; placenta; prostate; skeletal muscle; colon; male infertility; cancer; ss.
                                                                                                                                                                                                                                                                                                         28-JAN-2002; 2002EP-00001167.
                                                                                                                                                                                                                                                                                                                                       07-AUG-2002.
                                                                                                                                                                                                                                                                                                                                                                    EP1229046-A2
                                                                                                                                                                                                                               30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                    Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human HTPL scanning oligonucleotide
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                                                                                                                                                                                                              30-JAN-2001;
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                                                                                                                                    (AEOM-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25 GGAAGGCAAGCAG 37
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                                                                                                                                    AEOMICA
                                                                                                                                                              ; 2001WO-US000663.
; 2001WO-US000664.
; 2001WO-US000667.
; 2001WO-US000668.
; 2001WO-US000668.
; 2001WO-US000669.
; 2001US-00864761.
; 2001US-0327898P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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Pred. No.
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Example

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Page

91; 718pp; English

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Best Local
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30-JAN-2001;
30-JAN-2001;
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                                                                                                                                                                                                                                                                                                        07-AUG-2002
                                                                                                                                                                                                                                                                                                                                      EP1229046-A2
                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human HTPL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-JAN-2003
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 Novel isolated human testis for identifying agonist and
                                                                                                                                                                                                                                                                            28-JAN-2002;
                                               WPI; 2002-676582/73
                                                                                                                                                                                                                              30-JAN-2001;
                                                                                                          (AEOM-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        scanning oligonucleotide
                                                                                                                                   ; 2001WO-US000663.
; 2001WO-US000664.
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; 2001WO-US000669.
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expressed Patched like protein (HTPL), useful antagonist and specific binding partners, and
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RESULT 69
ABK19313/c
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Best Local
                                                                                                                                                                                                                                                                                                                                     Human; hammerhead ribozyme; cytostatic; antitumour; antidiabetic; ophthalmological; antitarthritic; antipsoriatic; virucide; osteopathic; vulnerary; cancer; lymphoma; Ewing's sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous sclerosis; port-wine stain; wound healing; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; ss; Osler-Weber-rendu syndrome, leukaemia; osteoporosis; DNAzyme; inozyme;
                                WPI; 2002-082995/11
                                                                Jarvis
                                                                                                                                                                                                                                                         WO200188124-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human ERG
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                                                                                                                                                   16-MAY-2000; 2000US-00572021
                                                                                                                                                                                   16-MAY-2001; 2001WO-US015866
                                                                                                                                                                                                                       22-NOV-2001.
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                                                                                                 RIBOZYME PHARM INC
GLAXO GROUP LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Amberzyme target sequence Seq'ID No 1960.
                                                                Von
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 5 A; 5 C; 6 G; 1 T; 0 U; 0 Other;
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                                                                Carlowitz I,
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Pred. No.
                                                                Mcswiggen
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2.5e+04;
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EJ
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Novel polynucleotide which down regulates expression of

Ets-related

gene

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RESULT 70
ABK19314/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity Matches 13; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                         tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous selerosis; port-wine stain; wound healing; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; 88;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 neovascular glaucoma, myopic degeneration, arthritis, psoriasis, verruca vulgaris, angiofibroma of tuberous sclerosis, port-wine stains, Sturge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          conditions selected from cancer, lymphoma, Ewing's sarcoma, mela tumour angiogenesis, diabetic retinopathy, macular degeneration,
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16-MAY-2000; 2000US-00572021
                                                                    16-MAY-2001; 2001WO-US015866
                                                                                                                                           22-NOV-2001
                                                                                                                                                                                                            WO200188124-A2
                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                    Osler-Weber-rendu syndrome,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ophthalmological; antiarthritic; antipsoriatic; virucide; osteopathic; vulnerary; cancer; lymphoma; Ewing's sarcoma; melanoma; psoriasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human ERG Amberzyme target sequence Seq ID No 1961.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hammerhead ribozyme; cytostatic; antitumour; antidiabetic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4.1%;
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                                                                                                                                                                                                                                                                                                                                                                                leukaemia; osteoporosis; DNAzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 13;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2.5e+04;
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tumour angiogenesis, diabetic retinopathy, macular degeneration, concerns angiogenesis, diabetic retinopathy, macular degeneration, concerns and move ascular glaucoma, myopic degeneration, arthritis, psoriasis, verruca concerns, angiofibroma of tuberous sclerosis, port-wine stains, Sturge conditions, port-wine stains, Sturge conditions, port-wine stains, Sturge concerns, port-wine stains, Sturge conditions, suitable for treatment, post-wine suitable for the treatment of ERG, conditions suitable for the treatment in conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or conjunction such as diagnost of conditions and conjunction such as diagnostic tool to the expression of ERG, and as diagnostic tool to conjunct of ERG RNA in a cell, (i) is useful for specifically conjuncted to the presence of ERG RNA in a cell, (i) is useful for specifically conjuncted to the presence of ERG RNA in a cell, (i) is useful for specifically conjuncted cardians and solidance of ERG, and specifically represent nucleic acids, including antisense and conjunction proposed that share homology with ERG gene or ERG fusions and conjunction proposed that share homology with ERG gene or ERG fusion genes.

Conjunction and conjunction of ERG, and
Sequence 17 BP; 4 A; 1 C; 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel polynucleotide which down regulates expression of Ets-related gene, useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           expression of an Ets-related gene (ERG). () conditions selected from cancer, lymphoma,
                                                            related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 4; Page 125; 149pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-082995/11.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GLAX ) GLAXO GROUP LTD
                                                         PCR primers of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       relates to
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G; 0 T; 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       acid molecule
U; 0 Other;
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                           Matches
                                Query Match
Best Local
       154 TTCCTTCCCCAGC 166
16
                           13;
                                 Similarity
                           Conservative
                                4.1%;
                           0
                                 Score 13;
Pred. No.
                           Mismatches
                                DB 6; L
                           ٥,
                                        Length 17;
                           Indels
                          0
                          Gaps
                           0
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RESULT 71
ABK19315/c
ID ABK19
09-APR-2002
                                                                Human
                                                                                  ABK19315;
                                                                                          ABK19315
                                                                                               2
                                                                ERG Amberzyme target sequence Seq ID No 1962.
                                                                                          standard;
                                                                         (first entry)
                                                                                          RNA;
                                                                                           17
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(RIBO-) RIBOZYME PHARM INC.

ophthalmological; antiarthritic; antipsoriatic; virucide; osteopathic; vulnerary; cancer; lymphoma; Ewing's sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; Human; hammerhead ribozyme; cytostatic; antitumour; antidiabetic; angiofibroma of tuberous sclerosis; port-wine stain; wound healing; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; leukaemia; DNAzyme; inozyme;

WO200188124-A2

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RESULT 72
ABV89388
ID ABV85
XX ABV85
XX ABV85
XX ABV85
XX ABV85
XX Human
XX Human
XX Human
XX Human
XX Gene
XX Gene
XX Gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CC Weber syndrome, Kippel-Trenauray-Weber syndrome, Osler-Weber-rendu (CC syndrome, leukaemia, osteoporosis and wound healing. (I) is useful for treating a patient having a condition associated with the level of ERG, by contacting cells of the patient with (I) under conditions suitable for the treatment. The method comprises the use of one or more therapies under conditions suitable for the treatment. Leukaemia or tumour angiogenesis is treated by administering (I) to the patient in conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or chemotherapy treatment. (I) is useful for reducing ERG activity in a cell, by contacting the cell with (I). (I) is useful for cleaving RNA of ERG gene, by contacting (I) with RNA, in the presence of a divalent cation such as Mg2+. (I) is useful for diagnosis of conditions and diseases related to the expression of ERG, and as diagnostic tool to examine genetic drift and mutations within diseased cells or to detect the presence of ERG RNA in a cell. (I) is useful for SRG fusion genes. ABK17354-ABK22719 represent nucleic acids, including antisense and carrenal are described by the cell such including antisense and carrenal are described by the cell such including antisense and carrenal are described by the cell such including antisense and carrenal are described by the cell such including antisense and carrenal are described by the cell such including antisense and carrenal are described by the cell such including antisense and carrenal are described by the cell such including antisense and carrenal are described by the cell such including antisense and carrenal are described by the cell such including antisense and carrenal are described by the cell such including antisense and carrenal are described by the cell such including antisense and carrenal are described by the cell such including antisense and carrenal are described by the cell such including antisense and carrenal are described by the cel
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 4 A;
EP1239051-A2
                                                                                                                 Human; POSHL 1; SH3 domain; POSH-like signalling Rho GTPase; signal transduction; gene expression;
                                                                                                                                                                                                                                               23-DEC-2002
                                                                                                                                                                                                                                                                                                                                            ABV89388 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 4; Page 125; 149pp; English.
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                                              Homo sapiens
                                                                                                                                                                                                 Human POSHL1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-MAY-2000; 2000US-00572021
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                                                                                            therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     154 TICCTICCCCAGC 166
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      primers of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.1%;
llarity 100.0%;
Conservative (
                                                                                                                                                                                           scanning oligonucleotide SEQ ID NO 101.
                                                                                                                                                                                                                                            (first entry)
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                                                                                            transgenic; ss.
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Pred. No. 2.5e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      G; 0 T; 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 17;
                                                                                                                   protein 1; oncogene;
; cancer; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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gene therapy;

Human; POSHL 1; SH3 domain; POSH-like signalling protein 1; oncogene; Rho GTPase; signal transduction; gene expression; cancer; vaccine;

Human POSHL1 scanning oligonucleotide

SEQ

ID NO

104.

ABV89391;

23-DEC-2002

(first entry)

ABV89391 standard; DNA; 17

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cc protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino CC (S1) having 95% deviations, especially conservative substitutions or a CC fragment of the sequences comprising at least 8 contiguous amino acids. CC Human POSHL 1 is a proto-oncogene/oncogene product that functions as an CC adaptor protein that interacts with Rho family small GTPases as well as CC downstream components of the signal transduction pathway. (I) is useful CC for identifying a specific binding partner. (I) and nucleic acids (II) CC caused by altered expression of human POSHL1 including diagnosing and CC treating cancer, they useful in the development of vaccines and (II) is useful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating CC are useful for measuring and for surveying gene expression and creating CC transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonuclectide useful in examples CC of the invention. Note: The present sequence did not form part of the CC Derwent by the European Patent Office
ABV89391
ID ABV89391
AZ ABV89391
AZ ABV89
XX ABV89
XX ABV89
XX ABV89
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Matches
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30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, -1, useful for treating disorders associated with decreased expressactivity of human POSHL1.
                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to an isolated SH3 domain protein 1 (POSHL 1) polypeptide (I), comprising
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23-MAY-2001;
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30-JAN-2001;
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                                                                                                                                                                                                                                                                                       l Similarity
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; 2001WO-US000667.
; 2001WO-US000669.
; 2001WO-US000669.
; 2001WO-US000670.
; 2001US-00864761.
                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                              BP;
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2001WO-US000665.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2002EP-00001165
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                                                                                                                                                                                                                                                                                                                                                C; 10
                                                                                                                                                                                                                                                                                                    Score 13;
Pred. No.
                                                                                                                                                                                                                                                                                                                                              G;
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                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                             T; 0
                                                                                                                                                                                                                                                                                                                                              U; 0 Other;
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hes 0;
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RESULT 74
ABV89392
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Best Local (
                                                                                                                                                                                                                                                                                                         Matches
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                 Human;
                                           Human POSHL1 scanning oligonucleotide SEQ ID NO 105.
                                                                                                                                                                                                                                                                                                                                                                                                 are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonuclectide useful in examples of the invention. Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to berwent by the European Patent Office
                                                                                  23-DEC-2002
                                                                                                                    ABV89392;
                                                                                                                                                   ABV89392 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                        Sequence 17
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               POSHL 1; SH3
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2001WO-US000666.
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                                                                             (first entry)
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domain; POSH-like signalling protein 1; oncogene;
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ive 0; Mismatches
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23-DEC-2002 ABV89390;

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The invention relates to an isolated SH3 domain (POSH)-like signalling CC protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino CC acids (SI, ABB3999), a sequence having 65% sequence identity to (SI), CC (SI) having 95% deviations, especially conservative substitutions or a creater of the sequences comprising at least 8 contiguous amino acids. CC Human POSHL 1 is a proto-oncogene/oncogene product that functions as an CC adaptor protein that interacts with Rho family small GTPases as well as CC downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (I) and nucleic acids (II) cencoding (I) are useful for diagnosing, monitoring disease and treating caused by altered expression of human POSHL1 including diagnosing and CC treating cancer, they useful in the development of vaccines and (II) is useful for measuring and for surveying gene expression and creating creater useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples of the invention. Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to Derwent by the European Patent Office
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Best Local Similarity
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30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000670.
                                                                                                                                                                                                                                                                            Sequence 17 BP; 2 A; 3 C; 11 G; 1 T; 0 U; 0 Other;
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GCGCGGAGGAGGG 13
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Human POSHL1 scanning oligonucleotide SEQ ID NO 103

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Matches 13; Conservative (
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Rho GTPase; signal transduction; gene expression; cancer; vaccine;
                                                                                                                                        Sequence 17 BP; 2 A; 3 C; 11 G; 1 T; 0 U; 0 Other;
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US-10-301-832-76
US-10-316-540-62
US-10-316-540-134
US-10-786-720-593
US-10-786-720-593
US-10-786-720-1609
US-10-786-720-2545
US-10-786-720-2545
US-10-786-720-2545
US-10-786-720-2545
US-10-786-720-2547
US-10-786-720-2547
US-10-786-720-2547
US-10-786-720-2549
US-10-786-720-2549
US-10-786-720-2549
US-10-060-756A-2210
US-10-060-756A-2211
US-10-060-756A-2211
US-10-060-756A-2211
US-10-080-756A-2211
US-10-080-756A-
                                                            ALIGNMENTS
                                                                                                                                                            Sequence 603, App Sequence 1042, App Sequence 1042, Appl Sequence 17, Appli Sequence 76, Appli Sequence 76, Appli Sequence 146, Appli Sequence 147, Appli Sequence 147, Appli Sequence 217, Appli Sequence 519, Appli Sequence 519, Appli Sequence 519, Appli Sequence 591, Appli Sequence 592, Appli Sequence 592, Appli Sequence 593, Appli Sequence 594, Appli Sequence 1606, Appli Sequence 1607, Appli Sequence 1607, Appli Sequence 1608, Appli Sequence 1609, Appli Sequence 2545, Appli Sequence 2546, Appli Sequence 2547, Appli Sequence 2548, Appli Sequence 2709, Appli Sequence 2709, Appli Sequence 2710, Appli Sequence 2711, Appli Sequence 2712, Appli Sequence 2713, Appli Sequence 2713, Appli Sequence 2713, Appli Sequence 2714, Appli Sequence 2715, Appli Sequence 2716, Appli Sequence 2717, Appli Sequence 2718, Appli Sequence 2718, Appli Sequence 2719, Appli Sequence 271
                                                                                           sequence
                                                                                             81420,
85086,
94187,
103434,
107646,
108888,
2163, A
2164, A
                                                                                                                                                                                                                                   ; LENGTH: 21
; TYPE: RNA
; ORGANISM: RNAi
US-10-751-736-15593
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Matches
  APPLICANT: Wyeth
APPLICANT: Marti
APPLICANT: Brown
APPLICANT: Liu,
TITLE OF INVENTIO
                                                                                                                                                                                       Matches
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Best Local &
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APPLICANT: Wyeth
APPLICANT: Myeth
APPLICANT: Brown, Eugene
APPLICANT: Liu, Wei
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TRE
TITLE OF INVENTION: CANCERS
PILE REFERENCE: AM100927 (031896-002000)
CURRENT APPLICATION NUMBER: US/10/751,736
CURRENT FILING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US Provisional Application 60/438,000
PRIOR FILING DATE: 2003-01-06
PRIOR FILING DATE: 2003-01-06
NUMBER OF SEQ ID NOS: 54873
SOPTWARE: Patentin version 3.2
SEQ ID NO 15593
LENGTH: 21
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APPLICANT: Brown, Eugene
APPLICANT: Brown, Eugene
APPLICANT: Liu, Wei
TITLE OF INVENTION: COMPOSITIONS AND METHODS
TITLE OF INVENTION: CANCERS
FILE REFERENCE: AM10927 (031896-002000)
CURRENT APPLICATION NUMBER: US/10/751,736
CURRENT FILING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US/10/751,736
CURRENT FILING DATE: 2003-01-06
PRIOR FILING DATE: 2003-01-06
NUMBER OF SEQ ID NOS: 54873
SOFTWARE: Patentin version 3.2
SEQ ID NO 15592
LENGTH: 21
                                                                                           RESULT 3
US-10-751-736-15685/c
US-10-751-736-15685/c
; Sequence 15685, Application US/10751736
; Publication No. US20040265230A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-10-751-736-15593/c
, Sequence 15593, Application US/10751736
, Publication No. US20040265230A1
, GENERAL INFORMATION:
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GENERAL INFORMATION:
APPLICANT: Wyeth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          146 TGCGACCTTTCCTTCC 161
                                                                                                                                                                                                                                     21 TGCGACCTTTCCTTCC
T: Brown, EugeneT: Liu, WeiINVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING
                                  Martinez, Robe
Brown, Eugene
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100.0%; Pr
                                                             Robert
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%; Pred. No. 2.6:
0; Mismatches
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Pred. No.
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2.6e+02;
hes 0;
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2.6e+02;
hes 0;
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RESULT 5
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; ORGANISM: homo sapiens
US-10-751-736-15685
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; ORGANISM: RNA1
US-10-751-736-15686
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Best Local S
Matches 16
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                                                                                                                                                                                                                                            Sequence 1167, Application US/09961077 Publication No. US20030014775A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO 15686
LENGTH: 21
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Best Local
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CURRENT APPLICATION NUMBER: US/10/751,736
CURRENT FILING DATE: 2003-01-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: AM100927 (031896-002000)
CURRENT APPLICATION NUMBER: US/10/751,736
CURRENT FILING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US Provisional Application 60/438,000
PRIOR FILING DATE: 2003-01-06
PRIOR FILING DATE: 2003-01-06
PRIOR FILING DATE: 2003-01-06
PRIOR FILING DATE: 2003-01-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US Provisional Application 60/438,000 PRIOR FILING DATE: 2003-01-06 NUMBER OF SEQ ID NOS: 54873
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                        TITLE OF INVENTION:
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ilarity 100.0%;
Conservative C
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Edington, Brent E.
McSwiggen, James A.
                                                                                         Young, Scott A.
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                                                        Folkerts, Otto
Merlo, Donald J.
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                                                                                                                                                 Patricia Ann Owens
 COMPOSITION AND METHODS FOR MODULATION OF GENE EXPRESSION IN PLANTS
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Pred. No.
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. 2.6e+02;
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US-09-961-077-1169
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                                                                                                                                                                                                                                                                                                  Sequence 1169, Application US/09961077 Publication No. US20030014775A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 4.7%; Score 15; Best Local Similarity 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 1167: SEQUENCE CHARACTERISTICS:
               Skokut, Thomas Skokut, Thomas Scott A.
Young, Scott A.
Young, Scott A.
Polkerte, Otto
Merlo, Donald J.
TITLE OF INVENTION: COMPOSITION AND METHODS FOR MODULATION OF GENE EXPRESSION
IN PLANTS
1263
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 08/679,645
FILING DATE: July 12, 1996
APPLICATION NUMBER: 60/001,135
FILING DATE: July 13, 1995
APPLICATION NUMBER: 08/300,726
FILING DATE: September 2, 1994
ATTORNEY,AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/09/961,077
FILING DATE: 21-Sep-2001
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. SOFTMARE: WOTCH PETECT 5.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 1263
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                  APPLICANT:
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TYPE: nucleic acid
STRANDEDNESS: single
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (213) 955-0440 TELEX: 67-3510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 219/247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Los Angeles
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McSwiggen, James A.
Merlo, Patricia Ann
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9.5e+02;
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Gaps

CITY: Los Angeles STATE: California COUNTRY: U.S.A.

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                                                                                    SOFTWARE: PatentIn version 3.2 SEQ ID NO 78 LENGTH: 18
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                                                                                                                                                                APPLICANT: Murphy, Finbarr
TITLE OF INVENTION: Apoptosis-Related Kinase/GPCRs
FILE REFERENCE: 8912/2012
CURRENT APPLICATION NUMBER: US/10/764,238
CURRENT FILING DATE: 2004-01-23
PRIOR APPLICATION NUMBER: US 60/457,533
PRIOR PILING DATE: 2003-03-25
PRIOR PILING DATE: 2003-03-25
PRIOR PILING DATE: 2003-01-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -10-764-238-78
                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: EiRx Therapeutics Ltd. APPLICANT: Seery, Liam
                                                                                                                                                       NUMBER OF SEQ ID NOS: 173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 1169:
SEQUENCE CHARACTERISTICS:
                         ORGANISM: Artificial sequence FEATURE:
                                                                TYPE: DNA
OTHER INFORMATION: QPCR Forward Primer (GPR86)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/679,645

FILING DATE: July 12, 1996

APPLICATION NUMBER: 60/001,135

FILING DATE: July 13, 1995

APPLICATION NUMBER: 08/300,726

FILING DATE: September 2, 1994

ATTORNEY/AGENT INFORMATION:

NAME: Warburg, Richard J.

REGISTRATION NUMBER: 32,327

REFERENCE/DOCKET NUMBER: 219/247

TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
les 15; Conserv
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COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear SEQ ID NO: 1169:
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OPERATING SYSTEM: IBM P.C.
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                   Seery, Liam
Hayes, Ian
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 21-Sep-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/09/961,077
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100.0%; Pred. No. 9.5e+02;
7ative 0; Mismatches 0;
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                                                                                                                                   ; ORGANISM: Homo sapien
US-10-098-263B-37841
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US-10-751-736-15688
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                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/098,263B
CURRENT FILING DATE: 2003-01-08
PRIOR APPLICATION NUMBER: 60/276,759
PRIOR FILING DATE: 2001-03-16
NUMBER OF SEQ ID NOS: 131066
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 37841
                                                                                                                                                                                                                                                                                                                    Sequence 37841, Application US/10098263B Publication No. US20030104410A1 GENERAL INFORMATION:
APPLICANT: Mittman, Michael TITLE OF INVENTION: Human Microarray FILE REFERENCE: 3118.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Liu, wei
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING
TITLE OF INVENTION: CANCERS
FILE REFERENCE: AM100927 (031896-002000)
CURRENT APPLICATION NUMBER: US/10/751,736
CURRENT FILING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US provisional Application 60/438,000
PRIOR PRIOR PRICE DATE: 2003-01-06
NUMBER OF SEQ ID NOS: 54873
SOFTWARE: Patentin version 3.2
SEQ ID NO 15688
LENGTH: 21
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Best Local Similarity
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Best Local &
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APPLICANT: Marti
APPLICANT: Brown
APPLICANT: Liu,
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                                                                                                                                                                     TYPE: DNA
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                                                                                                                                                                                     LENGTH: 25
                                176 GTCTCTGGAGATCAC 190
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17 GTCTCTGGAGATCAC 3
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15; Conserv
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Pred. No.
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US-10-296-012-13

; Sequence 13, Application US/10296012

; Publication No. US20040023238A1
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US-10-189-576-9
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US-09-740-332-118/c
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APPLICANT: Keller, Charles
APPLICANT: Ballard, Linda
Sequence 118, Application US/09740332
Publication No. US20030125270A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions
TITLE OF INVENTION: Hepatitis C Virus Infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 9, Application US/10189576
Publication No. US20030059878A1
GENERAL INFORMATION:
APPLICANT: OUUKI, Tetsuo
APPLICANT: KOGUCHI, Yutaka
APPLICANT: TSUBA, Naoki
TITLE OF INVENTION: A NOVEL PURINOCEPTOR AND THE GENE THEREOF
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Best Local (
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CURRENT APPLICATION NUMBER: US/10/189,576
CURRENT FILING DATE: 2002-07-02
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CURRENT APPLICATION NUMBER: US/10/296,012
CURRENT FILING DATE: 2003-06-23
PRIOR APPLICATION NUMBER: 60/219,531
PRIOR FILING DATE: 2000-07-20
NUMBER OF SEQ ID NOS: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn version 3.1 iEQ ID NO 9 LENGTH: 28
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EQ ID NO 13
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TITLE OF INVENTION: High Throughput Glutathione S-Transferase Polymorphic Allele Assa
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                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Artificially synthesized primer sequence
                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
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Lemons, Richard
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Conservative 0;
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100.0%; Pred. No. 9.4e+02;
ive 0; Mismatches 0;
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Pred. No.
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; OTHER INFORMATION: oligonucleotide substrate
US-09-740-332-4437
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US-09-740-332-4437
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FEATURE:
NAME/KEY: misc_feature;
LOCATION:
OTHER INFORMATION: oligonucleotide substrate
US-09-740-332-118
                                                                                                                                                                                                                                                                            RESULT 14
US-09-740-332-4438
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LENGTH: 17
TYPE: RNA
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LENGTH: 17
TYPE: RNA
                                                                                                                                                                                                               Sequence 4438, Application US/09740332 Publication No. US20030125270A1 GENERAL INFORMATION:
                               NUMBER OF SEQ ID NOS: 9704
SOFTWARE: PatentIn version
SEQ ID NO 4438
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Best Local Similarity
                                                                                           APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment
TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: RPI 400/003
CURRENT APPLICATION NUMBER: US/09/740,332
CURRENT FILING DATE: 2001-03-26
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TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relat
TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: RPI 400/003
CURRENT APPLICATION NUMBER: US/09/740,332
CURRENT FILING DATE: 2001-03-26
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CURRENT APPLICATION NUMBER: US/09/740,332
CURRENT APPLICATION DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9704
SOFTWARE: PatentIn version 3.0
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NUMBER OF SEQ ID NOS: 9704
SOFTWARE: Patentin version 3.0
LENGTH: 1'
TYPE: RNA
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Pred. No. 3.5e+03;
2; Mismatches 0
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ORGANISM: artificial sequence

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; OTHER INFORMATION: oligonucleotide substrate US-09-817-879-4437
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US-09-817-879-4437
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US-09-817-879-118/c
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APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: MBHB00-801-F
CURRENT APPLICATION NUMBER: US/09/817,879
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9703
SOUTWARE: PatentIn version 3.0
SEQ ID NO 118
LENGTH: 17
                                                                                                                                                                                                                     APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymetic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: MSHB00-801-F
CURRENT PAPPLICATION NUMBER: US/09/817,879
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9703
SOPTWARE: PatentIn version 3.0
SEQ ID NO 4437
       Best Local
Matches
                                           Query Match
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Best Local Similarity 85.7%;
Matches 12; Conservative
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Best Local Similarity
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Publication No. US20030171311A1
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                                                                                                                                PEATURE:
NAME/KEY: misc_feature
                                                                                                                                                                       ORGANISM: artificial sequence
                                                                                                                                                                                                 TYPE: RNA
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ORGANISM: artificial sequence
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     l Similarity 85.
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100.0%; Pred. No. 3.5e+03;
                   4.4%;
 Score 14; DB 10; Length 17; Pred. No. 3.5e+03; 2; Mismatches 0; Indels
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Pred. No. 3.5e+03;
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; TYPE: RNA; Homo sapiens US-10-712-672-98
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; OTHER INFORMATION: oligonucleotide substrate US-09-817-879-4438
                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/10/712/672
CURRENT FILING DATE: 2003-11-13
PRIOR APPLICATION NUMBER: US/09/653,225
PRIOR FILING DATE: 2000-08-31
PRIOR APPLICATION NUMBER: 60/197,769
PRIOR APPLICATION NUMBER: 60/197,769
PRIOR APPLICATION NUMBER: 60/150,713
PRIOR FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/150,713
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US-10-712-672-98/c
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SEQ ID NO 98
LENGTH: 17
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APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relat-
TITLE OF INVENTION: Hepatitis C Virus Infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 98, Applica Publication No. US20 GENERAL INFORMATION:
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CURRENT FILING DATE: 2001-03-26
NUMBER OF SEO ID NOS: 9703
SOFTWARE: PatentIn version 3.0
SEO ID NO 4438
LENGTH: 17
TYPE: RNA
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Best Local
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Best Local Similarity
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Publication No. US20030171311A1
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APPLICANT: Stinchcomb, Dan
TITLE OF INVENTION: Method and Reagent for the Inhibition of Telomerase Enzyme
FILE REFERENCE: MBHB00-882-C (400/019)
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                                                                                                           Similarity
                                        TCCTGAGGAAGCGG 118
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85.7*;
                                                                                   4.4%; Score 14; DB 17; 100.0%; Pred. No. 3.5e+03; tive 0; Mismatches 0;
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RESULT 19
US-10-712-672-99/c
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                                                                                                                                                            ; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-712-672-811
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US-10-712-672-811/c
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; ORGANISM: Homo sapiens
US-10-712-672-99
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Publication No. US20040102413A1
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Publication No. US20040102413A1
                                                                                                Query Match
Best Local Similarity
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APPLICANT: Stinchcomb, Dan
TITLE OF INVENTION: Method and Reagent for the Inhibition of Telomerase Enzyme
FILE REFERENCE: MBHB00-882-C (400/019)
CURRENT APPLICATION NUMBER: US/10/712,672
CURRENT FILING DATE: 2003-11-13
                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: US/09/653,225
PRIOR PILING DATE: 2000-08-31
PRIOR APPLICATION NUMBER: 60/197,769
PRIOR EILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/150,713
PRIOR FILING DATE: 1999-08-31
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PRIOR FILING DATE: 2000-08-31
PRIOR APPLICATION NUMBER: 60/197,769
PRIOR EILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/150,713
PRIOR EILING DATE: 1999-08-31
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14; Conservative 0;
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                                                                             Conservative 0;
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100.0%; Pred. No. 3.5
ive 0; Mismatches
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; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-712-672-812
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US-10-669-841-2711/c
                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: PCT/USO2/09187
PRIOR FILING DATE: 2002-03-26
PRIOR PELICATION NUMBER: US 60/296,876
PRIOR FILING DATE: 2001-06-08
PRIOR APPLICATION NUMBER: US 60/335,059
PRIOR APPLICATION NUMBER: US 60/337,055
PRIOR FILING DATE: 2001-10-24
PRIOR APPLICATION NUMBER: US 60/337,055
PRIOR FILING DATE: 2001-12-05
PRIOR FILING DATE: 2001-12-05
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APPLICANT: Kenneth, Draper
APPLICANT: Elisabeth, Roberts
TITLE OF INVENTION: OLIGONUCLEOTIDE MEDIATED INHIBITION OF HEPATITIS
TITLE OF INVENTION: VIRUS REPLICATION
FILE REFERENCE: 400/042US (MBHDG-249-E)
CURRENT APPLICATION NUMBER: US/10/659,841
CURRENT FILING DATE: 2003-09-23
CURRENT FILING DATE: 2003-09-23
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PRIOR FILING DATE: 2000-08-31
PRIOR APPLICATION NUMBER: 60/197,769
PRIOR FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/150,713
PRIOR FILING DATE: 1999-08-31
NUMBER OF SEQ ID NOS: 5586
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APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
TITLE OF INVENTION: Method and Reagent for the Inhibition of Telomerase Enzyme
FILE REFERENCE: MBHB00-882-C (400/019)
CURRENT APPLICATION NUMBER: US/10/712,672
CURRENT FILING DATE: 2003-11-13
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PRIOR FILING DATE: 2002-02-20
PRIOR APPLICATION NUMBER: US 60/363,124
PRIOR FILING DATE: 2002-03-11
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APPLICANT: Lawre
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                                  FILING DATE: 2001-03-26
APPLICATION NUMBER: US 09/740,332
                                                                                          APPLICATION NUMBER: US 09/817,879
APPLICATION NUMBER: US 09/611,931
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NUMBER OF SEQ ID NOS: 16207
SOFTWARE: PatentIn version 3.0
SEQ ID NO 2711
                                                                             SOFTWARE: PatentIn version SEQ ID NO 7030
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Matches 14; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/669,841
CURRENT FILING DATE: 2003-09-23
PRIOR APPLICATION NUMBER: PCT/US02/09187
PRIOR FILING DATE: 2002-03-26
PRIOR APPLICATION NUMBER: PCT/US02/09187
PRIOR FILING DATE: 2002-03-26
PRIOR APPLICATION NUMBER: PCT/US02/09187
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                                                                                                                      Remaining Prior Application data NUMBER OF SEQ ID NOS: 16207
                                                                                                                                                                 PRIOR FILING DATE: 2000-02-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 60/335,059 PRIOR FILING DATE: 2001-10-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: OLIGONUCLEOTIDE MEDIATED INHIBITION OF HEPATITIS B VIRUS AND HEPATITLE OF INVENTION: VIRUS REPLICATION FILE REFERENCE: 400/042US (MBHB02-249-E)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: US 09/504,321 PRIOR FILING DATE: 2000-02-15
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ORGANISM: Artificial Sequence FEATURE:
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OTHER INFORMATION: oligonucleotide substrate
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OTHER INFORMATION: Description of Artificial Sequence:
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                                                                                                                                                                                  APPLICATION NUMBER: US 09/504,321
                                                                                                                                                                                                       APPLICATION NUMBER: US 09/611,931 FILING DATE: 2000-07-07
                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 60/337,055
FILING DATE: 2001-12-05
                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 09/817,879
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Kenneth, Draper
Elisabeth, Roberts
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David, Morrissey
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                                                                                                                                                                                                                                                                                                                                                 ATION NUMBER: US 60/363,124
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PRIOR FILING DATE: 2002-03-26
PRIOR PPLICATION NUMBER: US 60/296,876
PRIOR APPLICATION NUMBER: US 60/335,059
PRIOR APPLICATION NUMBER: US 60/335,059
PRIOR FILING DATE: 2001-10-24
PRIOR APPLICATION NUMBER: US 60/337,055
PRIOR FILING DATE: 2001-12-05
PRIOR FILING DATE: 2001-12-05
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SEQ ID NO 7031
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Best Local &
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APPLICANT: Kenneth, Draper
APPLICANT: Eliasbeth, Roberts
TITLE OF INVENTION: OLIGONUCLEOTIDE MEDIATED INHIBITION OF HEPATITIS B VIRUS AND HEP?
TITLE OF INVENTION: VIRUS REPLICATION
FILE REFERENCE: 400/042US (MBHB02-249-E)
CURRENT APPLICATION NUMBER: US/10/669,841
CURRENT FILING DATE: 2003-09-23
CURRENT FILING DATE: 2003-09-23
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                                                                                                                                                                                                                                    TYPE: RNA
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                     ENGTH:
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APPLICATION NUMBER: US 60/363,124
APPLICATION SUMBER: US 60/363,124
APPLICATION NUMBER: US 60/363,124
                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US 09/740,332 FILING DATE: 2000-12-18
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David, Morrissey
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US-10-667-271-579/c
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Publication No.
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PRIOR FILING DATE: 2002-02-20
PRIOR APPLICATION NUMBER: USSN 60/363,124
PRIOR FILING DATE: 2002-03-11
PRIOR APPLICATION NUMBER: USSN 60/386,782
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CURRENT APPLICATION NUMBER: US/09/955,259B
CURRENT FILING DATE: 2002-08-30
                    PRIOR APPLICATION NUMBER: USSN 60/406,784
PRIOR FILING DATE: 2002-08-29
PRIOR APPLICATION NUMBER: USSN 60/408,378
PRIOR FILING DATE: 2002-09-05
PRIOR APPLICATION NUMBER: USSN 60/409,293
PRIOR FILING DATE: 2002-09-09
PRIOR FILING DATE: 2002-09-09
                                                                                                                                                                                                                                                                                                                  CURRENT FILING DATE: 2003-09-16
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR FILING DATE: 2003-05-23
PRIOR APPLICATION NUMBER: PCT / US03/05043
PRIOR PILING DATE: 2003-02-20
PRIOR PILING DATE: 2003-02-27
PRIOR APPLICATION NUMBER: PCT / US02/09187
PRIOR FILING DATE: 2002-03-26
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APPLICANT: MOTTISSEY, David
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Hepatitis C Virus (H
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (sina)
FILE REFERENCE: 400/129 (MBH802-763B)
CURRENT APPLICATION NUMBER: US/10/667,271
CURRENT FILING DATE: 2003-09-16
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SOFTWARE: PatentIn version 3.1
EQ ID NO 24
LENGTH: 19
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TITLE OF INVENTION: Expression of a Human Insulin
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PRIOR FILING DATE: 2002-08-05
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OTHER INFORMATION: Synthetic Primer
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Prior Application data
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No. US20040209831A1
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SEQ ID NO 580
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                                                                Matches
                                                                                               Query Match
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APPLICANT: Morrissey, David
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Hepatitis C Virus
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (sin
FILE REFERENCE: 400/129 (NBHB02-763B)
CURRENT APPLICATION NUMBER: US/10/667,271
CURRENT FILING DATE: 2003-09-16
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                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: USSN 60/409,293 PRIOR FILING DATE: 2002-09-09
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                                                                                                                                                 OTHER INFORMATION: Description of Artificial Sequence: Target
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                                                                                                                                                                                                        TYPE: RNA
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APPLICATION NUMBER: USSI
FILING DATE: 2002-09-05
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FILING DATE: 2002-03-11
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CURRENT FILING DATE: 2003-09-16
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR FILING DATE: 2003-05-23
PRIOR APPLICATION NUMBER: PCT / US03/05043
PRIOR APPLICATION NUMBER: PCT / US02/09187
PRIOR FILING DATE: 2003-02-26
PRIOR APPLICATION NUMBER: PCT / US02/09187
PRIOR APPLICATION NUMBER: USSN 60/401,104
PRIOR APPLICATION NUMBER: USSN 60/401,104
PRIOR APPLICATION NUMBER: USSN 60/401,104
PRIOR FILING DATE: 2002-08-05
                                                                                                                                                                                                                                                                 Sequence 592, Application US/10667271
Publication No. US20040209831A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics
APPLICANT: McSwiggen, James
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NUMBER OF SEQ ID NOS: 1705
SOFTWARE: PatentIn version 3.2
SEQ ID NO 584
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Best Local :
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                         APPLICANT: Macejāk, Dennis
APPLICANT: Beigelman, Leonid
APPLICANT: Beigelman, Leonid
APPLICANT: Morrissey, David
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Hepatitis C Virus
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (sin
FILE REPERENCE: 400/129 (MHHD02-763B)
CURRENT APPLICATION NUMBER: US/10/667,271
CURRENT FILING DATE: 2003-09-16
PRIOR APPLICATION NUMBER: US 10/444,853
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APPLICATION NUMBER: USSN 60/363,124
FILING DATE: 2002-03-11
APPLICATION NUMBER: USSN 60/386,782
FILING DATE: 2002-06-06
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FILING DATE: 2002-08-29
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Macejak, Dennis
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Macejak, Dennis
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                       2003-05-23
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PRIOR APPLICATION NUMBER: USSN 60/363,124
PRIOR FILING DATE: 2003-05-23
PRIOR APPLICATION NUMBER: PCT / US03/05043
PRIOR APPLICATION NUMBER: PCT / US03/05043
PRIOR APPLICATION NUMBER: PCT / US02/09187
PRIOR APPLICATION NUMBER: PCT / US02/09187
PRIOR APPLICATION NUMBER: USSN 60/401,104
PRIOR APPLICATION NUMBER: USSN 60/358,580
PRIOR APPLICATION NUMBER: USSN 60/363,124
PRIOR PILING DATE: 2002-02-20
PRIOR APPLICATION NUMBER: USSN 60/363,124
PRIOR FILING DATE: 2002-03-11
PRIOR APPLICATION NUMBER: USSN 60/366,782
PRIOR APPLICATION NUMBER: USSN 60/406,784
PRIOR APPLICATION NUMBER: USSN 60/406,784
PRIOR APPLICATION NUMBER: USSN 60/406,784
PRIOR APPLICATION NUMBER: USSN 60/408,378
PRIOR APPLICATION NUMBER: USSN 60/409,293
PRIOR FILING DATE: 2002-09-05
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US-10-667-271-592
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LENGTH: 19
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Best Local Similarity
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APPLICANT: Morrissey, David
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Hepatitis C Virus (HCV)
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (sina)
FILE REFERENCE: 400/129 (WBHB02-763B)
CURRENT APPLICATION NUMBER: US/10/667,271
CURRENT FILING DATE: 2003-09-16
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PRIOR APPLICATION NUMBER: USSN 60/409,293
PRIOR FILING DATE: 2002-09-09
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PRIOR APPLICATION NUMBER: USSN 60/406,784
PRIOR FILING DATE: 2002-08-29
PRIOR APPLICATION NUMBER: USSN 60/408,378
PRIOR APPLICATION NUMBER: USSN 60/408,378
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APPLICATION NUMBER: USSN 60/386,782
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APPLICATION NUMBER: USSN 60/358,580
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FILING DATE: 2002-03-26
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o. US20040209831A1
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File Wrapper
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CURRENT FILING DATE: 2003-09-16
PRIOR APPLICATION NUMBER: US/10/667,271
CURRENT FILING DATE: 2003-09-23
PRIOR APPLICATION NUMBER: PCT / US03/05043
PRIOR APPLICATION NUMBER: PCT / US03/05043
PRIOR FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT / US02/09187
PRIOR FILING DATE: 2002-03-26
PRIOR PILING DATE: 2002-03-26
PRIOR APPLICATION NUMBER: USSN 60/401,104
PRIOR FILING DATE: 2002-08-05
PRIOR FILING DATE: 2002-02-00
PRIOR FILING DATE: 2002-03-26
PRIOR FILING DATE: 2002-03-26
PRIOR FILING DATE: 2002-03-26
PRIOR FILING DATE: 2002-03-20
PRIOR APPLICATION NUMBER: USSN 60/363,124
PRIOR APPLICATION NUMBER: USSN 60/363,124
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US-10-667-271-620/c
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                                                                                                      Query Match
Best Local
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Best Local Similarity
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SEQ ID NO 597
                                                                                                                                                                                                                                                                                                                                             Remaining Prior Application data NUMBER OF SEQ ID NOS: 1705
                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: USSN 60/408,378 PRIOR FILING DATE: 2002-09-05
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PRIOR APPLICATION NUMBER: USSN 60/406,784
PRIOR FILING DATE: 2002-08-29
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                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn version 3.2
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                                                                                                                                                                                                                  FEATURE:
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                                                                                                                                                                                                                                                          LENGTH: 19
TYPE: RNA
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ORGANISM: Artificial Sequence
FEATURE:
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Macejak, Dennis
                                                                                  Conservative
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L00.0%; Pred. No.
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US-10-667-271-1275; Sequence, 1275; Application US/10667271; Publication No: US20040209831A1
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PRIOR FILING DATE: 2003-05-23
PRIOR PPLICATION NUMBER: PCT / US03/05043
PRIOR PPLICATION NUMBER: PCT / US02/09187
PRIOR APPLICATION NUMBER: PCT / US02/09187
PRIOR APPLICATION NUMBER: USSN 60/401,104
PRIOR APPLICATION NUMBER: USSN 60/458,580
PRIOR APPLICATION NUMBER: USSN 60/358,580
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Best Local S
Matches 12
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APPLICANT: McSwiggen, James
APPLICANT: Macejak, Dennis
APPLICANT: Beigelman, Leonid
APPLICANT: Morrissey, David
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                                      APPLICANT: Beigėlman, Leonid
APPLICANT: Morrissey, David
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Hepatitis C
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Aci
TILE REFERENCE: 400/129 (MBHB02-763B)
CURRENT APPLICATION NUMBER: US/10/667,271
CURRENT FILING DATE: 2003-09-16
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SOFTWARE: Patentin version 3.2
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PRIOR FILING DATE: 2002-09-05
PRIOR APPLICATION NUMBER: USSN 60/409,293
PRIOR FILING DATE: 2002-09-09
PRIOR APPLICATION NUMBER: US PRIOR FILING DATE: 2003-05-23
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CURRENT FILING DATE: 2003-09-16
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ORGANISM: Artificial Sequence
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Morrissey, David
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larity 85.7%;
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                    10/444,853
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; Pred. No. 3.5e
2; Mismatches
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APPLICATION NUMBER: PCT

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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: RNA Interference Mediated Inhibition of Hepatitis C Virus (HCV) TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (sina) FILE REFERENCE: 400/129 (MBHB02-763B) CURRENT APPLICATION NUMBER: US/10/667,271 CURRENT FILING DATE: 2003-09-16
                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR PILING DATE: 2003-05-23
PRIOR APPLICATION NUMBER: PCT / US03/05043
PRIOR FILING DATE: 2003-02-20
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APPLICANT: McSwi
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FILING DATE: 2002-03-26
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APPLICATION NUMBER: USSN 60/358,580
FILING DATE: 2002-02-20
                        APPLICATION NUMBER: USSN 60/409,293 FILING DATE: 2002-09-09
                                                                            FILING DATE: 2002-08-29
APPLICATION NUMBER: USSN 60/408,378
                                                                                                                    APPLICATION NUMBER: USSN 60/406,784
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APPLICATION NUMBER: USSN 60/386,782
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Macejak, Dennis
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Morrissey, David
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    Application data removed -
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TUMBER: PCT
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Pred. No. 3.56
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                                                                       ; OTHER INFORMATION: US-10-667-271-1288
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                                                                                                                                                         Remaining Prior Application data
NUMBER OF SEQ ID NOS: 1705
SOFTWARE: PatentIn version 3.2
SEQ ID NO 1288
LENGTH: 19
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SOFTWARE: PatentIn ver
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APPLICANT: Morrissey, David
APPLICANT: Morrissey, David
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Hepatitis C Virus (HCV)
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (sina)
FILE REFERENCE: 400/129 (MBHB02-763B)
CURRENT EPPLICATION NUMBER: US/10/667,271
CURRENT FILING DATE: 2003-09-16
FRIOR APPLICATION NUMBER: US 10/444,853
PRIOR APPLICATION NUMBER: PCT / US03/05043
PRIOR FILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-02-20
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PRIOR FILING DATE: 2002-03-26
PRIOR APPLICATION NUMBER: USSN 60/401,104
PRIOR FILING DATE: 2002-08-05
PRIOR APPLICATION NUMBER: USSN 60/358,580
PRIOR FILING DATE: 2002-02-20
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                                                                                                      TYPE: RNA
ORGANISM: Artificial
FEATURE:
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ORGANISM: Artificial
FEATURE:
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APPLICATION NUMBER: USSN 60/408,378
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FILING DATE: 2002-06-06
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Macejak, Dennis
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RESULT 37
US-10-667-271-1316
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US-10-667-271-1293
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PRIOR FILING DATE: 2003-05-23
PRIOR PPLICATION NUMBER: PCT / US03/05043
PRIOR PPLICATION NUMBER: PCT / US02/05187
PRIOR APPLICATION NUMBER: PCT / US02/09187
PRIOR APPLICATION NUMBER: USSN 60/401,104
PRIOR FILING DATE: 2002-08-05
PRIOR APPLICATION NUMBER: USSN 60/358,580
PRIOR APPLICATION NUMBER: USSN 60/363,124
PRIOR PILING DATE: 2002-02-02
PRIOR APPLICATION NUMBER: USSN 60/363,124
PRIOR PILING DATE: 2002-03-11
PRIOR APPLICATION NUMBER: USSN 60/386,782
PRIOR APPLICATION NUMBER: USSN 60/386,782
PRIOR APPLICATION NUMBER: USSN 60/386,782
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NUMBER OF SEQ ID NOS: 1705
SOFTWARE: PatentIn version 3.2
SEQ ID NO 1293
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APPLICANT: Beigelman, Leonid
APPLICANT: Morrissey, David
TITLE OF INVENTION: RAN Interference Mediated Inhibition of Hepatitis C Virus (HCV)
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (sina)
FILE REFERENCE: 400/129 (MBHB02-763B)
CURRENT APPLICATION NUMBER: US/10/667,271
CURRENT FILING DATE: 2003-09-16
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR FILING DATE: 2003-05-23
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APPLICANT: McSwiggen, James
APPLICANT: Macejak, Dennis
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APPLICANT: Morrissey, David
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Hepatitis C
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Aci
FILE REFERENCE: 400/129 (MBHB02-763B)
CURRENT APPLICATION NUMBER: US/10/667,271
CURRENT FILING DATE: 2003-09-16
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PRIOR APPLICATION NUMBER: USSN 60/406,784
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Macejak, Dennis
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US-10-028-248A-159
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CURRENT APPLICATION NUMBER: US/10/028,248A CURRENT FILING DATE: 2001-12-19
                                                  APPLICANT: Rothenberg, Mark
TITLE OF INVENTION: No. US20030235882Alel Nucleic Acids and Polypeptides and Methods
TITLE OF INVENTION: Thereof
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PRIOR FILING DATE: 2002-09-05
PRIOR APPLICATION NUMBER: USSN 60/409,293
PRIOR FILING DATE: 2002-09-09
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FILING DATE: 2002-08-05
APPLICATION NUMBER: USSN 60/358,580
FILING DATE: 2002-02-20
APPLICATION NUMBER: USSN 60/363,124
FILING DATE: 2002-03-11
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FILING DATE: 2002-06-06
APPLICATION NUMBER: USSN 60/406,784
FILING DATE: 2002-08-29
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FILING DATE: 2003-02-20
APPLICATION NUMBER: PCT / US02/09187
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Millet, Isabelle
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Miller, Charles
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Spytek, Kimberly
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Tchernev, Velizar
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Zerhusen, Bryan
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PRIOR APPLICATION NUMBER: 60/256619

2000-12-19

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RESULT 40
US-10-107-782-159
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                                                                                                                                                                                                                                                                                                                                                                                                                ; TYPE: DNA
; ORGANISM: Chlamydia pneumoniae
US-10-289-762-2554
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US-10-289-762-2554/c
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                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                            Sequence 159, Appropriate Publication No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Griffais, R.

TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence and polypeptides, fragments TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnosis, preve TITLE OF INVENTION: and treatment of infection FILE REFERENCE: 9710-003-999

CURRENT APPLICATION NUMBER: US/10/289,762

CURRENT FILING DATE: 2003-03-27

NUMBER OF SEQ ID NOS: 6849

SEQ ID NO 2554

LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 2554, Application US/10289762 Publication No. US20040006218A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 159
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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Best Local Similarity
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                                                                                                      APPLICANT: Boldog, Ferenc, APPLICANT: Casman, Stacie APPLICANT: Colman, Steve,
                                                     APPLICANT:
                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/285189
PRIOR FILING DATE: 2001-04-20
PRIOR APPLICATION NUMBER: 60/308039
PRIOR FILING DATE: 2001-07-26
PRIOR APPLICATION NUMBER: 60/311266
                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR FILING DATE: 2001-08-09
NUMBER OF SEQ ID NOS: 211
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/272408 PRIOR FILING DATE: 2001-02-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
OTHER INFORMATION: Description of Artificial Sequence:
OTHER INFORMATION: sequence
                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 2001-01-
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                                                                                                                                                                                                                                                                                                                 135
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                                                                    Casman, Stacie
Colman, Steve,
Edinger, Shlomit,
Gangolli, Esha,
                                                                                                                                                                                                                                                                               GGAGACGAGGTTGC 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CAGCACCAGTGGAA 20
Malyankar, Uriel,
                                                  Kekuda, Ramesh,
                                                                                                                                                                                          Application US/10107782
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                   Xiaohong,
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100.0%; Pred. No.
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RESULT 41
US-10-751-736-15496/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/10/107,782
CURRENT FILING DATE: 2002-03-27
PRIOR APPLICATION UNMBER: 10/028,248
PRIOR FILING DATE: 2001-12-19
PRIOR FILING DATE: 2001-12-6,619
PRIOR PILING DATE: 2000-12-19
PRIOR PILING DATE: 2000-12-19
PRIOR APPLICATION NUMBER: 60/262,959
PRIOR APPLICATION NUMBER: 60/272,408
PRIOR APPLICATION NUMBER: 60/272,408
PRIOR APPLICATION NUMBER: 60/285,189
PRIOR APPLICATION NUMBER: 60/285,189
PRIOR APPLICATION NUMBER: 60/285,189
                                                                                                                                                                                                                                                                                     Sequence 15496, Application US/10751736 Publication No. US20040265230A1 GENERAL INFORMATION:
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Matches
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SOFTWARE: CuraSeqList
SEQ ID NO 159
                                                                                                                                                                                            APPLICANT: Wyeth
APPLICANT: Martinez,
APPLICANT: Brown, Euc
APPLICANT: Liu, Wei
             TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING COLON TITLE OF INVENTION: CANCERS FILE REPERENCE; APAI00927 (031896-002000)

CURRENT APPLICATION NUMBER: US/10/751,736

CURRENT FILING DATE: 2003-01-06

PRIOR APPLICATION NUMBER: US Provisional Application 60/438,000

PRIOR APPLICATION NUMBER: US Provisional Application 60/438,000

RIOR APPLICATION NUMBER: US Provisional Application 60/438,000

RIOR FILING DATE: 2003-01-06
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PRIOR APPLICATION NUMBER: 60/279,344
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APPLICANT:
APPLICANT:
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PRIOR APPLICATION NUMBER: 60/308,039
PRIOR FILING DATE: 2001-07-26
PRIOR APPLICATION NUMBER: 60/311,266
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Zerhusen, Brian
TITLE OF INVENTION: NOVEL NUCLEIC ACIDS AND POLYPEPTIDES AND METHODS
FILE REFERENCE: 21402-225CIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
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mes 14; Conser
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PatentIn version 3.2
                                                                                                                                                                                                                Martinez, Robert
Brown, Eugene
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Tchernev, Velizar,
Vernet, Corine,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Smithson, Glennda,
Spytek, Kimberly,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sciore, Paul,
Shenoy, Suresh,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Stone, David,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Si, Jingsheng,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Shimkets, Richard
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Rothenberg, Mark,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Miller, Charles,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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)S: 215
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100.0%; Pred. No.
live 0; Mismatcl
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CURRENT APPLICATION NUMBER: US/09/883,152
CURRENT FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/211,835
PRIOR FILING DATE: 2000-06-15
NUMBER OF SEQ ID NOS: 127
SOFTWARE: FASESEQ for Windows Version 4.0
SEQ ID NO 53
LENGTH: 22
                                                                                                                                                                                                                                                                                                                              RESULT 43
US-10-098-263B-86601/c
| Sequence 86601, Application US/10098263B
| Publication No. US20030104410A1
| GENERAL INFORMATION:
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; TYPE: DNA
; ORGANISM: homo sapiens
US-10-751-736-15496
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                                                                                                  ; TYPE: DNA
; ORGANISM: Homo sapien
US-10-098-263B-86601
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                                 Query Match
Best Local S
Matches 14
                                                                                                                                                   NUMBER OF SEQ ID NOS: 131066
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 86601
LENGTH: 25
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Best Local Similarity 100.0%;
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APPLICANT: Kang, Sanmao
APPLICANT: Reinhard, Chri
                                                                                                                                                                                                                  APPLICANT: Mittman, Michael
TITLE OF INVENTION: Human Microarray
FILE REFERENCE: 3118.1
CURRENT APPLICATION NUMBER: US/10/098,263B
CURRENT FILING DATE: 2003-01-08
PRIOR APPLICATION NUMBER: 60/276,759
PRIOR FILING DATE: 2001-03-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Kang, Sanmao
APPLICANT: Reinhard, Christoph
APPLICANT: Jefferson, Anne Bennett
TITLE OF INVENTION: POLYNUCLEOTIDES RELATED TO COLON CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILE REFERENCE: 2300-1663
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ORGANISM: Artificial Sequence
FEATURE:
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                                 Local Similarity
mes 14; Conserv
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172 ACTGGTCTCTGGAG 185
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                             4.4%; Score 14; DB larity 100.0%; Pred. No. 3.: Conservative 0; Mismatches
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Pred. No.
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                                                 DB 15; 1
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                                                                                                                                                                     RESULT 46
                                                                                                Sequence 2463, Application US/10717597 Publication No. US20040110221A1 GENERAL INFORMATION:
                               APPLICANT: Wyeth
APPLICANT: Burcz
APPLICANT: Twine
APPLICANT: Dorne
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RESULT 44

US-10-098-263B-100111/c

US-10-098-263B-100111, Application US/10098263B

; Sequence 100111, Application US/10098263B

; Publication No. US20030104410A1

; GENERAL INFORMATION:
   APPLICANT: Mittman, Michael

; TITLE OF INVENTION: Human Microarray
                                                                                                                       ; LENGTH: 25
; TYPE: DNA
; ORGANISM: Homo sapien
US-10-098-263B-100737
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-10-098-263B-100737/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; ORGANISM: Homo sapien 
US-10-098-263B-100111
                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/276,759

PRIOR FILING DATE: 2001-03-16

NUMBER OF SEQ ID NOS: 131066

SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 100737
                                                                                                                                                                                                                                                                                                                                                                  Sequence 100737, Application US/10098263B
Publication No. US20030104410A1
GENERAL INFORMATION:
APPLICANT: Mittman, Michael
TITLE OF INVENTION: Human Microarray
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 131066
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 100111
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: 3118.1
CURRENT APPLICATION NUMBER: US/10/098,263B
CURRENT FILING DATE: 2003-01-08
PRIOR APPLICATION NUMBER: 60/276,759
PRIOR FILING DATE: 2001-03-16
                                                                                                                                                                                                                                                                                                             FILE REFERENCE: 3118.1
CURRENT APPLICATION NUMBER: US/10/098,263B
CURRENT FILING DATE: 2003-01-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
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172 ACTGGTCTCTGGAG 185
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                                        14;
                                                           Similarity
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                                        Conservative
                                                           4.4%; Score 14;
100.0%; Pred. No.
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100.0%; Pred. No.
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                                        Mismatches
                                                           DB 15;
. 3.5e+03;
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Gaps

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Slonim, Donna K. Twine, Natalie C. Dorner, Andrew J. Trepicchio, William L. Burczynski, Michael E.

Gaps

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CURRENT APPLICATION NUMBER: US/10/717,597
CURRENT FILING DATE: 2003-11-21
PRIOR APPLICATION NUMBER: US 60/459,782
PRIOR EILING DATE: 2003-04-03
PRIOR APPLICATION NUMBER: US 60/427,982
PRIOR EILING DATE: 2002-11-21
NUMBER OF SEQ ID NOS: 4904
SOFTWARE: PatentIn version 3.2
SEQ ID NO 2463
LENGTH: 25
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; ORGANISM: Homo sapiens
US-10-717-597-2463
                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: Oligonucleotide primers US-09-960-244A-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Wagner, Joseph
TITLE OF INVENTION: CELL POPULATIONS WHICH CO-EXPRESS CD49c
TITLE OF INVENTION: AND CD90
FILE REFERENCE: 2831.2003-000
CURRENT PAPLICATION NUMBER: US/09/960,244A
CURRENT FILING DATE: 2001-09-21
NUMBER OF SEQ ID NOS: 16
SOFTWARE: FeatSEQ for Windows Version 4.0
SEQ ID NO 11
LENGTH: 26
                                                                                                             GENERAL INFORMATION:
APPLICANT: Ho, Tony W.
APPLICANT: Kopen, Gen
                                                                                                                                                                    Sequence 11, Application US/10251685 Publication No. US20040058412A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 11, Application US/09960244A
Publication No. US20030059414A1
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APPLICANT: Kopen, Gene C.
APPLICANT: Righter, Willi
APPLICANT: Rutkowski, J.
APPLICANT: Wagner, Joseph
                                                                             APPLICANT:
       TITLE OF
                                         APPLICANT:
                                                            APPLICANT:
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TITLE OF INVENTION: METHODS FOR DIAGNOSING RCC AND OTHER SOLID TUMORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: AM101080L
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nes 14; Conserv
                                                                                                                                                                                                                                                                                                             105 TCCTGAGGAAGCGG 118
T: Herring, W. Joseph
T: Ragalia, Vanessa
'INVENTION: CELL POPULATIONS WHICH CO-EXPRESS CD49c
                                                                                                                                                                                                                                                                             24
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                                                                   Kopen, Gene C.
Righter, William F.
Rutkowski, J. Lynn
                                                        Wagner, Joseph
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                                                                                                                                                                                                                                                                                                                                                     Conservative
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100.0%; Pr
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100.0%; Pred. No. 3.5e+03;
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RESULT 50
US-10-257-017B-96936/c
; Sequence 96936, Application US/10257017B
; Publication NO. US20040241651A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; NUMBER OF SEQ ID NOS: 16
; SOFTWARE: FastSEQ for Windows Ve
; SEQ ID NO 11
; LENGTH: 26
; TYPE: DNA
; ORGANISM: Artificial Sequence
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US-10-257-017B-96935
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Best Local Similarity
PRIOR APPLICATION NUMBER: DE 10019173.8 PRIOR FILING DATE: 2000-04-07 NUMBER OF SEQ ID NOS: 382046
                                                                                TITLE OF INVENTION: Detection of single nucleotide polymorhphisms [SNPs] and cytosin TITLE OF INVENTION: methylations FILE REPERENCE: E01/1193/WO CURRENT APPLICATION NUMBER: US/10/257,017B CURRENT FILING DATE: 2002-10-07
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                                                                                                                                                                            APPLICANT: Alexander Olek
APPLICANT: Christian Piepenbrock
APPLICANT: Kurt Berlin
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APPLICANT: Christian Piepenbrock
APPLICANT: Kurt Berlin
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CURRENT FILING DATE: 2002-09-20
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FILE REFERENCE: 2831.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE: OTHER INFORMATION: Oligonucleotide for detection of SNP TSC0024052
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
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les 13; Conserv
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Pred. No.
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Pred. No.
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, OTHER INFORMATION: Oligonuclectide for detection of SNP TSC0024052 US-10-257-017B-96936
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US-10-257-017B-195240
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                                                                                                                                                                                                                                                                                                                                           Sequence 195240, Application US/10257017B Publication No. US20040241651A1 GENERAL INFORMATION:
                                                                                        NUMBER OF SEQ ID NOS: 382046
SEQ ID NO 195240
LENGTH: 13
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                                                                                                                                                                                    TITLE OF INVENTION: Detection of single nucleotide polymorhphisms [SNPs] and cytosine TITLE OF INVENTION: methylations FILE REFERENCE: E01/1193/WO CURRENT APPLICATION NUMBER: US/10/257,017B CURRENT FILING DATE: 2002-10-07
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APPLICANT: Christian Pier
APPLICANT: Kurt Berlin
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APPLICANT: Christian Piel
                                                                                                                                                PRIOR APPLICATION NUMBER: DE 10019173.8 PRIOR FILING DATE: 2000-04-07
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ORGANISM: Artificial Sequence
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      FEATURE: OTHER INFORMATION: Oligonucleotide for detection of SNP TSC0048037
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                                                      ORGANISM: Artificial Sequence
                                                                            TYPE: DNA
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Kurt Berlin
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Kurt Berlin
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ilarity 100.0%;
Conservative (
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ilarity 100.0%;
Conservative
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;; Pred. No. 1.3
0; Mismatches
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Pred. No.
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. 1.3e+04;
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APPLICANT: Macejak, Dennis
TITLE OF INVENTION: ENZYMATIC NUCLEIC ACID TREATMENT OF
TITLE OF INVENTION: ENZYMATIC NUCLEIC ACID TREATMENT OF
TITLE OF INVENTION: HERATITIS C VIRUS INFECTION
FILE REFERENCE: rpi 247/282
CURRENT APPLICATION NUMBER: US/09/504,231A
CURRENT FILING DATE: 2000-02-15
PRIOR APPLICATION NUMBER: 09/274,553
PRIOR FILING DATE: 1999-03-23
PRIOR FILING DATE: 1999-03-23
PRIOR APPLICATION NUMBER: 09/257,608
PRIOR APPLICATION NUMBER: 09/267
PRIOR APPLICATION NUMBER: 09/083,217
PRIOR PILING DATE: 1998-09-18
PRIOR APPLICATION NUMBER: 09/083,217
PRIOR FILING DATE: 1998-09-18
PRIOR PILING DATE: 1998-09-18
PRIOR PILING DATE: 1998-09-18
PRIOR PILING DATE: 1998-09-18
PRIOR PILING DATE: 1998-09-18
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                                                              US-09-504-231A-212
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; ORGANISM: Homo sapiens
US-10-416-928-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patent No. US...
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LENGTH: 14
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                                                                                                                                                                                        SEQ ID NO 212
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                     Query Match
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CURRENT APPLICATION NUMBER: US/10/416,928
CURRENT FILING DATE: 2003-05-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: METHOD FOR ANALYZING A PATIENT'S PREDISPOSITION TO INSULIN-DEPEN TITLE OF INVENTION: DIABETES, DEVICE AND SET OF PRIMERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 14
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                                                                                                                                                                                                             SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                  OTHER INFORMATION: Description of Artificial Sequence: Nucleic Acid Target
                                                                                                     ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                 TYPE: RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
                                                                                                                                                                  LENGTH: 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       193 TAGAATTGTAGTG 205
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Roberts, Beth
Pavco, Pamela
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4.1%; Score 13; 100.0%; Pred. No.
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;; Pred. No.
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hes 0;
                     DB 9;
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                     Length 15
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Matches

Local Similarity

13; Conservative

Mismatches

1.3e+04;

Indels

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Gaps

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, OTHER INFORMATION: Description of Artificial Sequence:capture
US-10-297-068-671
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US-09-274-553D-212/c
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Best Local S
                                                                                                                   SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 671
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 671, Application US/10297068 Publication No. US20030228585A1
                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 3148
SOFTWARE: PatentIn version 3.0
SEQ ID NO 212
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                                                                                                                                                               FILE REFERENCE: 13140P1174
CURRENT APPLICATION NUMBER: US/10/297,068
CURRENT FILING DATE: 2002-11-27
PRIOR APPLICATION NUMBER: JP 2000-164798
PRIOR FILING DATE: 2000-06-01
NUMBER OF SEQ ID NOS: 1298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Macejak, Dennis
TITLE OF INVENTION: ENZYMATIC NUCLEIC ACID TREATMENT OF DISEASES OR CONDITIONS RELATI
TITLE OF INVENTION: HEPATITIS C VIRUS INFECTION
FILE REFERENCE: 101 247/282
CURRENT APPLICATION NUMBER: US/09/274,553D
CURRENT FILING DATE: 1999-032
CURRENT FILING DATE: 1999-032
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PRIOR FILLING DATE: 1999-02-24
PRIOR APPLICATION NUMBER: 60/100,842
PRIOR FILLING DATE: 1998-09-18
PRIOR APPLICATION NUMBER: 60/083,217
PRIOR FILLING DATE: 1998-04-27
                                                                                                                                                                                                                                                                                       APPLICANT: MORIYA, Shogo
APPLICANT: MISHIDA, Michio
TITLE OF INVENTION: KIT AND METHOD FOR DETERMINING HLA TYPES
                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: INOKO, Hidetoshi
APPLICANT: KAGIYA, Taeko
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                     TYPE: DNA
                                                           ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     129 TCGGGCGGAGACG 141
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US20020082225A1
                                                                                                                                                                                                                                                                                                                                                         ICHIHARA, Tatsuo
Matsumura, Yoshiyuki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                McSwiggen,
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hes 0;
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                                                                                                           ; OTHER INFORMATION: Description of Artificial Sequence: Nucleic Acid US-09-864-785-2831
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US-09-864-785-2831/c
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                                    Query Match
Best Local Similarity
Matches 13; Conserv
                                                                                                                                                                                                      SOFTWARE: PatentIn version 3.0
SEQ ID NO 2831
LENGTH: 17
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 2831, Application US/09864785
Patent No. US20020177568A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 2830, Application US/09864785 Patent No. US20020177568A1
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                                                                                                                                                                                                                                                                             APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Enzymatic Nucleic Acid
TITLE OF INVENTION: Levels of NF-Kappa B
FILE REFERENCE: 400/022 (MBHB00-812-D)
CURRENT APPLICATION NUMBER: US/09/864,785
CURRENT FILING DATE: 2001-05-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Stinchcomb, Dan
APPLICANT: Draper, Ken
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Enzymatic Nucleic Acid
TITLE OF INVENTION: Levels of NF-Kappa B
FILE REFERENCE: 400/022 (MBHB00-812-D)
                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 3929
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/864,785
CURRENT FILING DATE: 2001-05-23
NUMBER OF SEQ ID NOS: 3929
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                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                FEATURE:
                                                                                                                                                                 ORGANISM: Artificial Sequence
                                                                                                                                                                                      TYPE: RNA
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 154
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TTCCTTCCCCAGC 166
                                                                                                                                                                                                                                                                                                                                                                                         Draper, Ken
                                                                                                                                                                                                                                                                                                                                                                                                              Stinchcomb, Dan
                                  4.1%; Score 13; DB 9; L
llarity 100.0%; Pred. No. 1.3e+04;
Conservative 0; Mismatches 0;
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; Pred. No. 1.3e+04
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Pred. No.
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                                                                     Length 17
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FILE REFERENCE: 400/022 (MBHB00-812-D)
CURRENT APPLICATION NUMBER: US/09/864,785
CURRENT FILING DATE: 2001-05-23
NUMBER OF SEQ ID NOS: 3929
SOFTWARB: PatentIn version 3.0
SEQ ID NO 2832
LENGTH: 17
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US-09-740-332-117/c
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US-09-864-785-2832/c
                                                                                                                                                                                                                                                                                US-09-740-332-117
                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 9704
SOFTWARE: PatentIn version 3.0
SEQ ID NO 117
LENGTH: 17
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Draper, Ken
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases
TITLE OF INVENTION: Levels of NF-Kappa B
TITLE OF INVENTION: Levels of NF-Kappa B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals Inc.

TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions

TITLE OF INVENTION: Hepatitis C Virus Infection

FILE REFERENCE: RPI 400/003

CURRENT APPLICATION UNMEER: US/09/740,332

CURRENT FILING DATE: 2001-03-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 117, Application US/09740332 Publication No. US20030125270A1
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 APPLICANT:
                                                                                                                                                                                                                                                                                                                          NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                               TYPE: RNA
ORGANISM: artificial sequence
FEATURE:
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                                                                                                                                                                                                                                                                                           OTHER INFORMATION: oligonucleotide substrate
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les 13; Conserv
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Ribozyme Pharmaceuticals Inc.
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100.0%; Pred. No. 1.
ive 0; Mismatches
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                                                                                                                                                              US-10-060-756A-210
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LENGTH: 17
TYPE: RNA
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                                                                                                                                                                           SEQ ID NO 210
LENGTH: 17
TYPE: DNA
ORGANISM: Homo sapiens
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GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 13; Conserv
                                                                                                                      Query Match
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PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9703
SOFTWARE: PatentIn version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
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TITLE OF INVENTION: HUMAN TESTIS EXPRESSED PATCHED LIKE PROTEIN
FILE REFERENCE: PB0177
                                                                                                                                                                                                                                                               SOFTWARE: Aeomica Sequence Listing Engine
                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 2001-10-09
NUMBER OF SEQ ID NOS: 4804
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NAME/KEY: misc_feature
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                                                                                               Local
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APPLICATION NUMBER: US 09/864,761
FILING DATE: 2001-05-23
APPLICATION NUMBER: US 60/327,898
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GGAAGGCAAGCAG
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                                                                              4.1%; Score 13; DB 14; larity 100.0%; Pred. No. 1.3e+04 Conservative 0; Mismatches 0
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US-10-060-756A-211
; Sequence 211, Application US/10060756A
; Publication No. US20030046717A1

RESULT 63

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US-10-060-756A-212; Sequence 212, App.; Publication No. U
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-060-756A-212
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                                                                                         SOFTWARE: Aec
SEQ ID NO 212
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PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
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                                                                                                                                                                   PRIOR FILING DATE:
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TITLE OF INVENTION: HUMAN TESTIS EXPRESSED PATCHED LIKE PROTEIN
FILE REFERENCE: PB0177
FILE REFERENCE: PB0177
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TITLE OF INVENTION: HUMAN TESTIS EXPRESSED PATCHED LIKE PROTEIN
FILE REFERENCE: PB0177
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ORGANISM: Homo sapiens
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00665
                                                                                                                                                                                     FILING DATE: 2001-05-23
APPLICATION NUMBER: US 60/327,898
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13; Conserv
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o. US20030046717A1
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                                                                                                               Sequence Listing Engine
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                                                     PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR PFLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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SEQ ID NO 213
LENGTH: 17
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Publication No. US20
GENERAL INFORMATION:
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Best Local Similarity
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TITLE OF INVENTION: HUMAN TESTIS EXPRESSED PATCHED LIKE PROTEIN
FILE REFERENCE: PB0177
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
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CURRENT FILING DATE: 2002-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
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NUMBER OF SEQ ID NOS: 4804
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PRIOR APPLICATION NUMBER: US 60/327,898
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PRIOR FILING DATE: 2001-01-30
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PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 09/864,761
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                     APPLICATION NUMBER: PCT/US01/00668 FILING DATE: 2001-01-30
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o. US20030046717A1
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o. US20030046717A1
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100.0%; Pred. No.
PCT/US01/00663
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Pred. No.
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RESULT 68
US-10-061-201-102
; Sequence 102, Application US/10061201
; Publication No. US20030166229A1
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-060-756A-214
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TITLE OF INVENTION: HUMAN POSH-LIKE PROTEIN
FILE REFERENCE: PB0178
FULL APPLICATION NUMBER: US/10/061,201
CURRENT FILING DATE: 2002-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00665
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Best Local Similarity
Matches 13; Conserv
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Best Local Similarity
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PRIOR APPLICATION NUMBER: US 09/864,761
PRIOR FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/327,898
PRIOR FILING DATE: 2001-10-09
NUMBER OF SEQ ID NOS: 4804
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PRIOR FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/328,205
PRIOR FILING DATE: 2001-10-10
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Homo sapiens
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b. US20030166229A1
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llarity 100.0%; Pred. No. 1.3e+0.
Conservative 0; Mismatches
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US-10-061-201-103
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-061-201-102
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PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR TILING DATE: 2001-01-30
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CURRENT FILING DATE: 2002-01-30
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TITLE OF INVENTION: HUMAN POSH-LIKE PROTEIN
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NUMBER OF SEQ ID NOS: 4162
SOFTWARE: Aeomica Sequence
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PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
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PRIOR APPLICATION NUMBER: PCT/US01/00669
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CURRENT FILING DATE: 2002-01-30
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FILING DATE: 2001-01-30
APPLICATION NUMBER: US 09/864,761
FILING DATE: 2001-05-23
                   APPLICATION NUMBER: US 09/864,761 FILING DATE: 2001-05-23
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00663
FILING DATE: 2001-01-30
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Similarity 100.0%; Pred. No.
13; Conservative 0; Mismatc
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o. US20030166229A1
NUMBER: US 60/328,205
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; ORGANISM: Homo sapiens
US-10-061-201-103
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US-10-061-201-104
                 Sequence 105, Application US/10061201
Publication No. US20030166229A1
GENERAL INFORMATION:
APPLICANT: Shannon, Mark
TITLE OF INVENTION: HUMAN POSH-LIKE PROTEIN 1
FILE REFERENCE: PB0178
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SEQ ID NO 104
LENGTH: 17
TYPE: DNA
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SOFTWARE: Acomica Sequence Listing
SEQ ID NO 103
LENGTH: 17
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CURRENT FILING DATE: 2002-01-30
CURRENT APPLICATION NUMBER: US/10/061,201
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NUMBER OF SEQ ID NOS: 4162
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TITLE OF INVENTION: HUMAN POSH-LIKE PROTEIN 1
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100.0%; Pred. No. 1.3e+04
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TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions;
FILE REFERENCE: MBH800-876-N (400/049)
CURRENT APPLICATION NUMBER: US/10/138,674;
CURRENT FILING DATE: 2002-05-03;
NUMBER OF SEQ ID NOS: 20822;
SOFTWARE: Patentin version 3.0
SEQ ID NO 7348
LENGTH: 17
Type: """
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; ORGANISM: Homo sapiens
US-10-061-201-105
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US-10-138-674-7348
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                                                                   Query Match 4.3
Best Local Similarity 84.0
Matches 11; Conservative
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Best Local Similarity
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PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
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PRIOR APPLICATION NUMBER: US 60/328,205
PRIOR FILING DATE: 2001-10-10
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PRIOR APPLICATION NUMBER: US 09/864,761
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
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PRIOR APPLICATION NUMBER: PCT/US01/00664
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APPLICATION NUMBER: PCT/US01/00670
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1.3e+04;
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RESULT 73 US-10-287-949A-7348

7348, Application US/10287949A

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RESULT 75
US-10-712-672-810/c
IS-10-712-672-810/c
Sequence 810, Application US/10712672
Publication No. US20040102413A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Chowrira, Bharat
APPLICANT: McSwiggen, Jim
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; ORGANISM: Homo sapiens
US-10-712-672-100
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US-10-712-672-100/c
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CURRENT FILING DATE: 2003-04-11
NUMBER OF SEQ ID NOS: 20822
SOFTWARE: Patentin version 3.0
SEQ ID NO 7248
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APPLICANT: Stinchcomb, Dan
TITLE OF INVENTION: Method and Reagent for the Inhibition of Telomerase Enzyme
FILE REFERENCE: MBHB00-882-C (400/019)
CURRENT APPLICATION NUMBER: US/10/712,672
CURRENT FILING DATE: 2003-11-13
CURRENT FILING DATE: 2003-11-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: US/09/653,225
PRIOR FILING DATE: 2000-08-31
PRIOR APPLICATION NUMBER: 60/197,769
PRIOR FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/150,713
PRIOR FILING DATE: 1999-08-31
NUMBER OF SEQ ID NOS: 5586
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TYPE: RNA
ORGANISM: Homo sapiens
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Local Similarity 84.6%;
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McSwiggen, Jim
Stinchcomb, Dan
Escobedo, Jaime
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APPLICANT: Stinchcomb, Dan
TITLE OF INVENTION: Method and Reagent for the Inhibition of Telomerase Enzyme
FILE REFERENCE: MBHB00-882-C (400/019)
CURRENT APPLICATION NUMBER: US/10/712,672
CURRENT FILING DATE: 2003-11-13
PRIOR APPLICATION NUMBER: US/9/653,225
PRIOR PILING DATE: 2000-08-31
PRIOR PILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/197,769
PRIOR FILING DATE: 2000-04-14
PRIOR FILING DATE: 2000-04-14
PRIOR FILING DATE: 1999-08-31
NUMBER OF SEQ ID NOS: 5586
SOFTWARE: Patentin version 3.0
SEQ ID NO 810
LENGTH: 17
TYPE: RNA
ORGANISM: Homo sapiens
US-10-712-672-810

Query Match
Best Local Similarity 100.0%; Pred. No. 1.3e+04;
Matches 13; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Matches 13; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Search completed: February 3, 2005, 00:11:21 Job time: 177.942 secs

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US-09-18-452A-2554
US-09-18-256BA-190
US-09-064-115A-190
US-09-018-595B-7
US-09-324-709A-7
US-09-324-709A-7
US-09-344-521-40
US-09-484-521-40
US-09-255-2028-773
US-08-240-547-11
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US-08-240-547-11
US-08-240-547-11
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US-08-863-639A-95
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US-08-913-635-13
US-08-954-210-11
US-08-918-937-6550
US-08-918-934-210-11
US-09-311-7728-5550
US-08-909-742-3
US-08-909-742-3
US-08-909-742-4
US-08-998-099-13
US-08-998-099-13
US-08-998-099-13
US-08-998-099-13
US-09-412-289-3
US-09-313-4199-1
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US-09-866-108A-8276
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US-09-866-108A-8277
US-09-9866-108A-8277
US-09-9866-108A-8277
US-09-9866-108A-8277
US-09-970-740-4
US-09-108-1163
US-09-108-6948-1163
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RESULT 1
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                                                                                                                                                                  Sequence 1, Application US/08651692
Patent No. 5856099
                                                                                                  GENERAL INFORMATION:
APPLICANT: Loren Miraglia, Thomas Geiger,
APPLICANT: Clarence Frank Bennett and Nicholas M. Dean
TITLE OF INVENTION: Compositions and Methods for
TITLE OF INVENTION: Modulating Type I Interleukin-1 Rev
                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 210 Lake Driv
CITY: Cherry Hill
STATE: NJ
COUNTRY: USA
ZIP: 08002
COMPUTER READABLE FORM:
                                                                                 NUMBER OF SEQUENCES: 4
                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                              E: Law Offices
210 Lake Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                 of Jane Massey Licata
East, Suite 201
                                                                                                                                                                                                                                                                                                           US-09-866-108A-13165
US-09-866-108A-13166
US-09-866-108A-13168
US-09-866-108A-13170
US-09-866-108A-13171
US-09-866-108A-13172
US-09-866-108A-13173
US-09-866-108A-13173
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US-09-866-108A-13174
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US-09-866-108A-13176
US-09-866-108A-13176
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US-09-422-978-9566
US-09-422-978-9566
US-09-122-624-13
US-09-182-762-6
US-09-181-780-15
US-09-177-650-15
US-09-177-650-65
US-08-570-155-16
US-08-570-155-16
US-08-570-155-16
US-08-570-155-17
US-08-23-300A-23
US-09-318-138-42
US-09-318-138-42
US-09-318-138-42
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US-08-374-144-3
US-08-775-164-3
US-08-775-609-3
US-08-775-607-3
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US-09-259-411-8
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US-09-742-693-1
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US-08-863-639A-67
                                                                                                                                                                                                                                         ALIGNMENTS
                                                                                                       Type I Interleukin-1 Receptor Expression
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RESULT 2
US-08-679-645-1167
; Sequence 1167, Application US/08679645
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Patent No. 635(
GENERAL INFORM APPLICANT:
APPLICANT:
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STRANDEDNESS: Single
; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-08-651-692-1
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REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH
TELECOMMUNICATION INFORMATION:
TELEPHONE: (609) 779-2400
TELEPAX: (609) 779-8488
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 16
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MEDIUM TYPE: DISKETTE, 3.5
COMPUTER: IBM PS/2
OPERATING SYSTEM: PC-DOS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION UNMER: US/08/6
FILING DATE: Herewith
                                                                                                                                                                                                                                                                                                                                 STREET: Suite 4700
STREET: Suite 4700
CITY: Los Angeles
CTATE: California
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TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                 COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C.
SOFTWARE: WOR'D PERfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth
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ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: US/08/679,645 FILING DATE: July 12, 1996 CLASSIFICATION: 800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH:
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Edington, Brent E.
McSwiggen, James A.
Merlo, Patricia Ann C
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M PS/2
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PRIOR APPLICATION DATA:

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Best Local Similarity
Tarkches 15; Conserva
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                                                                                               COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5" Diskette, 1.44 Mb

MEDIUM TYPE: storage

COMPUTER: IBM Compatible

OPERATING SYSTEM: IBM P.C. DOS 5.0

SOFTWARE: Word Perfect 5.1

CURRENT APPLICATION DATA:

APPLICATION UMBER: US/08/679,645

FILING DATE: Us/12, 1996

CLASSIFICATION: 800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: September 2, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 219/247
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Folkerts,
APPLICANT: Merlo, Don
TITLE OF INVENTION: C
TITLE OF INVENTION: M
TITLE OF INVENTION: I
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   guence 1169, Ap
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/001,135
FILING DATE: July 13, 1995
APPLICATION NUMBER: 08/300,726
FILING DATE: September 2, 1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                              STREET: 633 West Fi
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
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FILING DATE: JULY 13, 1995
APPLICATION NUMBER: 08/300,726
FILING DATE: September 2, 1994
                                                                                                                                                                                                                                                                           COUNTRY:
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nucleic acid
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633 West Fifth Street
Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Edington, Brent E. McSwiggen, James A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATION:
Zwick, Michael G.
Brent E.
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Merlo, Donald J.
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100.0%; Pred. No.
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                                                                                                       US-08-863-639A-27
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                                                   Matches
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Best Local Similarity
                                                                         Query Match
                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: JOSEPH E. Mueth
20,532
REGISTRATION UNMBER: 20,532
REFERENCE/DOCKET NUMBER: 1185
TELECOMMUNICATION INFORMATION:
TELEPHONE: (626) 795-6321
TELEFAX: (626) 795-6321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 21
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 24 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Coassin, Peter J.
APPLICANT: Rampal, Jang B.
APPLICANT: Caskey, C. T.
TITLE OF INVENTION: OLIGONUCLECTIDE REPEAT ARRAYS
NUMBER OF SEQUENCES: 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS
LENGTH: 18 base pairs
                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
                                                                                                                     MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 222
STREET: 222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 1011
TELEFAX: 67-3510
                                                                 Local
                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: Corel WordPerfect 8 version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: FILING DATE: May 28 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE:
                                                                                                                                 TOPOLOGY:
                                                                                                                                           STRANDEDNESS:
                                                                                                                                                            TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Sheldon & Mak
STREET: 225 South Lake Avenue, 9th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27, 5981185
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                                                                                                                                                          H: 24 base pairs
nucleic acid
                                                                 Similarity
                       ceccecceccaec 53
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                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matson, Robert S.
Coassin, Peter J.
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                                                   Conservative
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                                                                                                                                   linear
                                                                                                                                                                                                                                                                                                            May 28, 1997
DN: 435
                                                                                                                                                                                                                                                                                                                                                                                                        Diskette,
                                                                                                                     Other nucleic acid
                                                                                                                                             single
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                                               100.0%; --
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100.0%; Pr
                                                                                                                                                                                                                                                                                                                                   US/08/863,639A
                                                                                                                                                                                                                                                                                                                                                                                                        3.50 inch, 1.44 Mb storage
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                                                                Score 15; pred. No.
                                                                                                                                                                                                                                                       11859-1
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                                                   Mismatches
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o. 2.5e+02;
0;
                                                                 2.6e+02;
                                                                            Length 24;
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US-08-651-692-2
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                                  SEQ ID NO 15
LENGTH: 20
                                                                                                                                                                                                  Sequence 15, Application US/09513729B Patent No. 6165791 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 2, Application US/08651692
Patent No. 5856099
                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                           APPLICANT: Ian Popoff
APPLICANT: Jacqueline Wyatt
TITLE OF INVENTION: ANT
                                                                        CURRENT APPLICATION NUMBER: US/09/513,729B
CURRENT FILING DATE: 2000-02-24
NUMBER OF SEQ ID NOS: 88
                                                                                                                               FILE REFERENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (609) 779-8488 INFORMATION FOR SEQ ID NO:
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Loren Miraglia, Thomas Geiger,
APPLICANT: Clarence Frank Bennett and Nicholas M. Dean
TITLE OF INVENTION: Compositions and Methods for
TITLE OF INVENTION: Modulating Type I Interleukin-1 Receptor Expression
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Jane Massey Licata
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: IS!
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08,
FILING DATE: Herewith
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PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 210 Lake L
                                                                                                                                                                                                                                                                                                                                                                                                                              Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM F5/2
OPERATING SYSTEM: PC-DOS
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                                                                                                                                                                                                                                                                                                                                    14 GCCGCCGCAGCCCT 1
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14; Conservative (
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                                                                      2000-02-24
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                                                                                                                                                 ANTISENSE MODULATION OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ISPH-0144
                                                                                                                                                                                                                                                                                                                                                                                                                          Score 14; DB 2;
; Pred. No. 8.6e+0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; ORGANISM: Chlamydia pneumoniae
US-09-198-452A-2554
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-182-968A-190/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Griffals, R.
TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence and polypeptides, fra-
TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnosis,
TITLE OF INVENTION: and treatment of infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 6849
SEQ ID NO 2554
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequence 2554, Application US/09198452A
Patent No. 6559294
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 190, Application US/08182968A Patent No. 5610054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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Best Local Similarity
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Best Local &
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                         COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.
MEDIUM TYPE: 8torage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/182,
FILING DATE: 13-JANUARY-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/882,888
                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE: 9710-003-999
CURRENT APPLICATION NUMBER: US/09/198,452A
                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Draper, I
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH:
FILING DATE: 14-MAY-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                          CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                            ADDRESSEE: Lyon & Lyon STREET: 633 West Fifth Street STREET: Suite 4700
                                                                                                                                                                                                                        COUNTRY: U.S.A. ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                135 GGAGACGAGGTTGC 148
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100.0%; Pr
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                                                                                                                                                                                                                                                                                                                                                                                Kenneth G.
METHOD AND REAGENT FOR INHIBITING HEPATITIS C VIRUS REPLICATION
                                                                                                                                      IBM P.C. DOS 5.0
                                  07/882,888
                                                                                  US/08/182,968A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 14;
; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 14; DB 3; L
Pred. No. 8.6e+02;
                                                                                                                                                                                          1.44 Mb
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RESULT 9
US-08-774-306A-190/c
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                                                 US-08-774-306A-190
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Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/774,300
FILLING DATE: December 26, 1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/182,968
FILLING DATE: January 13, 1994
APPLICATION NUMBER: 07/882,888
FILLING DATE: May 14, 1992
ATTORNEY/ACENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (213) 955-044
TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                                                                          INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Draper, FITTLE OF INVENTION: TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER REALABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
                                                                                                                                                                                   NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 223/227
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: IBM P.C. DOS 5.0 SOFTWARE: Word Perfect 5.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                 ELECUTE: (213)
                                                                                                                                               TELEFAX: 1211.
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                                                              STRANDEDNESS: 811
TOPOLOGY: linear
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TOPOLOGY: lir
                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: U.S.A. ZIP: 90071-2066
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    r: Suite 4700
Los Angeles
                                                                                           nucleic acid
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633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Draper, Kenneth G.
VENTION: METHOD AND REAGENT FOR
VENTION: INHIBITING HEPATITIS C
VENTION: VIRUS REPLICATION
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4.1%;
100.0%;
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; Pred. No.
Score 13;
Pred. No.
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o. 2.9e+03;
0;
DB 2; L
2.9e+03;
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              Length 15;
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RESULT 11
US-09-018-595B-7
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; TOPOLOGY: lii
US-09-064-156A-190
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                                                                                                                                     Matches
                                                                                                                                                     Query Match
Best Local S
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TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/064,1
FILING DATE: April 21, 1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/774,306
APPLICATION NUMBER: 08/774,306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Draper, K
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: May 14, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: 23
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 08/774,306
FILING DATE: December 26, 1996
APPLICATION NUMBER: 08/182,968
FILING DATE: January 13, 1994
APPLICATION NUMBER: 07/882,888
                                                                                                                                                                                                                                                  TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                         TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM:
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STREET: Suite 4700
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                                                                                       129 TCGGGCGGAGACG 141
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                                                                                                                                  h 4.1%; Score 13; DB 3; Length 15;
Similarity 100.0%; Pred. No. 2.9e+03;
13; Conservative 0; Mismatches 0; Indels
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METHOD AND REAGENT FOR
INHIBITING HEPATITIS C
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Sequence 7, Application US/09018595B
Patent No. 5962233
GENERAL INFORMATION:
APPLICANT: Perkin-Elmer Corporati
APPLICANT: Applied Biosystems Div

Perkin-Elmer Corporation, Applied Biosystems Division

TITLE OF INVENTION:

DETERMINATION OF GENOTYPE OF AULTIPLE ALLELIC SITES

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US-09-324-709A-7
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                                                                                                                                                                                                                                                                                                 Sequence 7, Application US/09324709A
Patent No. 6154707
GENERAL INFORMATION:
APPLICANT: Perkin-Elmer Corporation,
APPLICANT: Applied Biosystems Division
TITLE OF INVENTION: FLUORESCENCE GENOTYPING AT MULTIPLE ALLELIC
TITLE OF INVENTION: SITES
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                        ZIP: 94304-1050
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: David J. Weitz
REGISTRATION NUMBER: 38,362
REFERENCE/DOCKET NUMBER: PELM-744
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 493-9300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM compatible
OPERATING SYSTEM: Microsoft Windows 95
SOFTWARE: Wordpetfect for windows 6.0,
SOFTWARE: ASCII (DOS) TEXT format
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/018,595B
                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: David J. Weitz,
ADDRESSEE: Wilson Sonsini Goodrich & Rosati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 94304-1050
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch diskette
                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
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COMPUTER: IBM compatible
OPERATING SYSTEM: Microsoft Windows 95
SOFTWARE: Wordperfect for windows 6.0,
SOFTWARE: ASCII (DOS) TEXT format
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                                                                                                                                                                                            STREET: 650 Page Mill Road CITY: Palo Alto
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TOPOLOGY: linear
                                                                                                                                                      COUNTRY:
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STATE: California
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                                                                                                                                                                        California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                David J. Weitz,
Wilson Sonsini Goodrich & Rosati
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APPLICANT: Harris Busch
APPLICANT: Jacqueline Wyatt
TITLE OF INVENTION: ANTISENSE MODULATION OF GU PROFILE REFERENCE: RTS-0163
CURRENT APPLICATION NUMBER: US/09/844,521
CURRENT FILING DATE: 2001-04-27
NUMBER OF SEQ ID NOS: 87
SEQ ID NO 40
LENGTH: 20
TYPE: DNA
                                              Sequence 10, Application US/09438906
Patent No. 6465185
GENERAL INFORMATION:
APPLICANT: Goldfine, Ira
APPLICANT: Trischitte, Vincenzo
APPLICANT: Pizzuti, Antonio
APPLICANT: Vigneri, Riccardo
APPLICANT: Frittitta, Lucia
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                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Thehes 13; Conserve
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                                                                                                                                                                          US-09-438-906-10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 40, Application US/09844521
Patent No. 6492172
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                APPLICANT: Frittitta, Lucia
TITLE OF INVENTION: Polymorphic Human PC-1 Sequences
FILE REFERENCE: 9076-089
CURRENT APPLICATION NUMBER: US/09/438,906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: C. Frank Bennett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (650) 493-6811
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 nucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
ATTORY AGENT INFORMATION:
ANAME: David J. Weitz
NAME: David J. Weitz
REGISTRATION NUMBER: 38,362
REFERENCE/DOCKET NUMBER: 1684
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 493-9300
                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: sing
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US-08-240-547-21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; TYPE: DNA; ORGANISM: H. US-09-438-906-10
                                                                                                                               US-08-240-547-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                              Matches
                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT FILING DATE: 1999-11-18
PRIOR APPLICATION NUMBER: 60/108,853
PRIOR FILING DATE: 1998-11-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: FastSEQ for Windows Version SEQ ID NO 10 LENGTH: 22
                                                                                                                                                                                                                                           TELEFAX: (510) 814-2977 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Sias Ph.D., Stacey R.
REGISTRATION NUMBER: 32,630
REFERENCE/DOCKET NUMBER: 8586
TELECOMMUNICATION INFORMATION:
                                                                                                                                           TOPOLOGY: 11
                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 23 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/918,844
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Resnick, Robert M.
APPLICANT: Young, Karen K.Y.
TITLE OF INVENTION: Primers and Probes for Detection of
TITLE OF INVENTION: Hepatitis C and No. 5527669el Variants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: U.S.A. ZIP: 07110-1199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: 340 F
CITY: Nutley
                                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/240,547
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                      140 CGAGGTTGCGACC 152
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                                                             l Similarity
13; Conserv
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CGAGGTTGCGACC 13
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340 Kingsland Street
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                                                              Conservative
                                                                                                                                                                                                                                                                            (510) 814-2863
                                                                                                                                            DNA (genomic)
                                                                                                                                                                           single
                                                           4.1%; Score 13; DB
100.0%; Pred. No. 2.9
tive 0; Mismatches
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                                                                            DB 1; Lo
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RESULT 17
US-09-225-928-773
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US-08-859-998-773
                                                                                                                                                                            Sequence 773, Applicat. Patent No. 6352829
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Field, Bret E.
REGISTRATION NUMBER: 37,620
REFERENCE/DOCKET NUMBER: 0909
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-322-5070
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US
PILING DATE: 21-MAY-19
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
COMPUTER: IBM Compat
OPERATING SYSTEM: Wi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Chenchik, Alex
APPLICANT: Jokhadze, George
APPLICANT: Bibilashvilli, Robert
TITLE OF INVENTION: METHOD OF ASSAYING DIFFERENTIAL
TITLE OF INVENTION: EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: FastSEQ for Windows Version CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: DNA FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 94025
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH:
                CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 2200 Sand Hill Road, Suite 100
                                                                                      Bibilashvilli, Robert
TITLE OF INVENTION: METHOD OF ASSAYING DIFFERENTIAL
EXPRESSION
                                                                                                                                     APPLICANT: Chenchik, Alex
Jokhadze, George
                                                                   NUMBER OF SEQUENCES: 1375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 773, Application US/08859998
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CITY :.. Menlo Park
                                                                                                                                                                                                                                                                                                   CGGCTCCTGCGTC 19
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2200 Sand Hill Road, Suite 100
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SYSTEM: Windows95
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; Pred. No. 2.9e+C
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ATTORNEY/AGENT INFORMATION:

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US-09-225--
; Sequence 773, Pr.
; Sequence 773, Pr.
; GENERAL INFORMATION:
APPLICANT: Chenchik, Alex
Jokhadze, George
Dibilashvilli, Rob
TYPRESS!
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OTHER INFORMATION: oligonucleotide primer;
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SEQUENCE DESCRIPTION: SEQ ID NO: 773;
US-09-225-928-773
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US-09-225-201B-773
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REGISTRATION NUMBER: 37,620
REFERENCE/DOCKET NUMBER: 0909.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-322-5070
TELEPHONE: 415-854-0875
INFORMATION FOR SEQ ID NO: 773:
SEQUENCE CHARACTERISTICS:
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Best Local Similarity
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COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: PastSEQ for Windows Version 2.0
CURRENT APPLICATION NUMBER: US/09/225,201B
FILING DATE: 05-Jan-1999
CLASSIFICATION : -Unknown>
PRIOR APPLICATION NUMBER: US/08/859,998
FILING DATE: 21-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Diskette
COMPUTER: LBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FRAUSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: METHOD OF ASSAYING DIFFERENTIAL EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 08/859,998 FILING DATE: 21-MAY-1997 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/09/225,928
FILING DATE: 05-Jan-1999
CLASSIFEICATION: «Unknown»
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: line MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
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TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                            STATE: CA
COUNTRY: US
                                                                                                                                                                                                                                                                                                  CITY: Menlo Park
                                                                                                                                                                                                                                                                                                              ADDRESSEE: Fish & Richardson, P.C. STREET: 2200 Sand Hill Road, Suite 100
                                                                                                                                                                                                                                        ZIP: 94025
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ZIP: 94025
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APPLICATION

PILING DATE:

ATTORNEY/AGENT INFORMATION:

NAME: Sias Ph.D., Stacey R.

REGISTRATION NUMBER: 32,630

REFERENCE/DOCKET NUMBER: 8586

TELEPHONE: (510) 814-2863

TELEPHONE: (510) 814-2977

INFORMATION FOR SEQ ID NO: 14:

SEQUENCE CHARACTERISTICS:
LENGTH: 24 base pairs

TYPE: mucleic acid

TYPE: mucleic acid
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US-08-240-547-14/c
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TELEFAX: 415-854-0875
INFORMATION FOR SEQ LID NO: 773:
SEQUENCE CHARACTERISTICS:
LENGTH: 23 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: Resnick, Robert M.
APPLICANT: Young, Karen K.Y.
TITLE OF INVENTION: Primers and Probes for Detection of
TITLE OF INVENTION: Hepatitis C and No. 5527669el Variar
NUMBER OF SEQUENCES: 43
                                                                                                                                                                                                                                                                CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/918,844
                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDLUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMAPPLICATION DATA:
APPLICATION NUMBER: US/08/240,547
TOPOLOGY: lir
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Nutley
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: U.S.A. ZIP: 07110-1199
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Field, Bret E.
REGISTRATION NUMBER: 37,620
REFERENCE/DOCKET NUMBER: 09096/002001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    E: Hoffmann-La Roche Inc.
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                    linea
DNA (genomic)
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Pred. No.
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2.9e+03;
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APPLICANT: Tonai, Richard
APPLICANT: StemCyte, Inc.
TITLE OF INVENTION: High Throughput Methods of HLA Typing
FILE REFERENCE: 020035-00210US
CURRENT APPLICATION NUMBER: US/09/747,391
CURRENT FILING DATE: 2001-07-13
PRIOR APPLICATION NUMBER: US 60/172,768
PRIOR FILING DATE: 1999-12-20
NUMBER OF SEQ ID NOS: 278
SOFTWARE: FastSEQ for Windows Version 3.0
ENGTH: 30
TUPE: 30
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                                                            Matches
                                                                                         Query Match
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                                                                            Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                        Patent No. 66701
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -09-344-529-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/344,529
CURRENT FILING DATE: 1999-06-24
EARLIER APPLICATION NUMBER: US 60/090,794
EARLIER FILING DATE: 1998-06-26
EARLIER APPLICATION NUMBER: US 60/095,713
EARLIER FILING DATE: 1998-08-07
NUMBER OF SEQ ID NOS: 19
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APPLICANT: HSC Research and Development Limited Partnership
TITLE OF INVENTION: Sculpin-Type Antifreeze Polypeptides and Nucleic Acids
FILE REFERENCE: 016252-002620US
FILE REFERENCE: 016252-002620US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                     TYPE: DNA ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Description of Artificial Sequence:upper strand OTHER INFORMATION: PCR primer
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ORGANISM: Artificial Sequence
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                                                                            100.0%;
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100.0%; Pred. No.
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2.9e+03;
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. 2.9e+03;
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US-08-609-657-10
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                                                                                                                                                                                                                            Patent No.
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Matches 12; Conserva
                                                                                                                                                                                                                              Sequence 10, Application US/08609657 Patent No. 5681706
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                                                                                                                                                                                                               GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                               APPLICANT:
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APPLICANT: Coassin, Peter J.
APPLICANT: Rampal, Jang B.
APPLICANT: Caskey, C. T.
COMPUTER READABLE FORM:
                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Hodgson, Russ,
                                                                                                                                  APPLICANT: Stoler, Daniel L. TITLE OF INVENTION: Mammalia TITLE OF INVENTION: Element
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ADDRESSEE: Sheldon & Mak
                                                                                                                       NUMBER OF SEQUENCES:
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                                              STATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Joseph E. Mueth
REGISTRATION NUMBER: 20,
REFERENCE/DOCKET NUMBER:
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SOFTWARE: Corel WordPerfect 8 version
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STREET: 225 South Lake Avenue, 9th Floor
                                                         CITY: Buffalo
                                                                        STREET:
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                            COUNTRY:
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                           United States
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                                                                                                                                                                                                 Anderson, Garth R.
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N: 435
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100.0%; Pred. No.
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9.8e+03;
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; HYPOTHETICAL:
US-08-609-657-10
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US-08-985-162-1759
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Best Local Similarity
                                                                                                                                          APPLICATION NUMBER: US/08/985,162
FILING DATE: 04 December 1997
CLLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/036,476
FILING DATE: 31 January 1997
ATTORNEY/AGENT INFORMATION:
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INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERITICS:
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TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
UNMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: ISSUE COMPUTER: COMPUTER:
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CURRENT APPLICATION DATA:
APPLICATION UNBER: US/08/600 CT
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 230/107
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FastSEQ for Windows 2.0
CURRENT APPLICATION DATA:
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NAME: Nelson, M. Bud
NAME: Nelson, M. Bud
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Akhtar, Saghir
APPLICANT: Fell, Patricia
APPLICANT: McSwiggen, James
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TELECOMMUNICATION INFORMATION:
TELLEPHONE: (716) 856-4000
TELEPAX: (716) 849-0349
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US-09-401-063-1759
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Matches
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GENERAL INFORMATION:
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TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                                                             INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                  SOFTWARE: FASTSEN: IBM P.C. DOS 5.0 SOFTWARE: FASTSEQ for Windows 2.0 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/401 062 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
APPLICANT:
                                            SEQUENCE CHARACTERISTICS
LENGTH: 14 base pairs
TYPE: nucleic acid
                                                                                                                                      NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 236
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/985,162

FILING DATE: 04 December 1997

APPLICATION NUMBER: 60/036,476

FILING DATE: 31 January 1997
                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Lyon & Ly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
             STRANDEDNESS: si
                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM CONTROL OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                         TELEPHONE: (21
TELEFAX: (213)
                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: U.S.A.
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TOPOLOGY: lir
                                                                                                               TELEX:
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McSwiggen, James
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                                                                                                                                                                                                                                                                                                                                                                                                          IBM Compatible
                               single
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OF DISEASES OR CONDITIONS RELATED
TO LEVELS OF EPIDERMAL GROWTH
FACTOR RECEPTORS
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; Pred. No. 9.8e+03;
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Query Match Best Local Similarity

3.8%; Score 12; 100.0%; Pred. No.

DB 4; Length 14; 9.8e+03;

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RESULT 26
US-08-863-639A-21/c
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                                                                                                                                                                                                                                                                                                                                Query Match 3.8%; Score 12; Best Local Similarity 100.0%; Pred. No. Matches 12; Conservative 0; Mismatc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21, Application US/08863639A Patent No. 5981185
                                                                                                                                          Sequence 1, Application US/08618834C Patent No. 6361937 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: JOSEPH E. MUSCH
REGISTRATION NUMBER: 20,532
REFERENCE/DOCKET NUMBER: 11859-1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (626) 796-4000
                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: Other nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM: Windows 95
SOFTWARE: Corel Wordberfect 8 version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/863,639A
FILING DATE: May 28, 1997
CLASSIFICATION: 435
                 NUMBER OF SEQUENCES: 54
CORRESPONDENCE ADDRESS:
ADDRESSEE: Ritter, Van Pelt & Yi LLP
STREET: 4906 El Camino Real, Suite 205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                      APPLICANT: Stryer, Lubert
TITLE OF INVENTION: Computer-Aided Nucleic Acid
TITLE OF INVENTION: Sequencing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Sheldon &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Matson, Robert S.
APPLICANT: Coassin, Peter J.
APPLICANT: Rampal, Jang B.
APPLICANT: Caskey, C. T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (626)
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225 South Lake Avenue, 9th Floor
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795-6321
NO: 21:
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US-08-618-834C-7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequence 7, Application US/08618834C Patent No. 6361937
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APPLICANT: Stryer, Lubert
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                                                                                                                                      SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: 105/08/618,834C FILLING DATE: 19-MAR-1996 CLASSIFICATION: 435 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
NAME: Ritter, Michael J.
REGISTRATION NUMBER: 36,653
REFERENCE/DOCKET NUMBER: AF
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-903-3500
                                                                                       ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: 650-903-3500
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ATTORNEY/AGENT INFORMATION:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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CITY: Los Altos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Ritter, Michael J.
REGISTRATION NUMBER: 36,653
REFERENCE/DOCKET NUMBER: AF
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OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                         APPLICATION NUMBER:
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4906 El Camino Real, Suite 205
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RESULT 30
US-08-954-210-11/c
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US-08-730-635-13/c
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                                                                                                                                                                                     Matches
                                                                                                                                                                                                                        Query Match
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NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 3.8%; Score 12; DB 3; I Best Local Similarity 100.0%; Pred. No. 9.8e+03 Matches 12; Conservative 0; Mismatches 0.
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                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/730,635
FILING DATE: 11-OCT-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Bak, Mary E.
REGISTRATION NUMBER: 31,215
REFERENCE/DOCKET NUMBER: B&FTUSA
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 540-9200
TELEPAX: (215) 540-5818
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Lansdo:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/730,635
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Lansdorp, Peter
TITLE OF INVENTION: Method for Detecting Multiple Copies of
TITLE OF INVENTION: a Repeat Sequence in a Nucleic Acid Molecule
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                           MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (2)
                                                                                                                                                                                   Local Similarity 100.0%;
nes 12; Conservative (
                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid STRANDEDNESS: sing TOPOLOGY: linear
                                                                                                                                                                                                                                                                                              TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
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                                                                                                           CGCCGCCGCCGC 2
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321 No. 6514693ristown Road
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                                                                                                                                                                                                   3.8%; Score 12; DB 4; Length 15; 100.0%; Pred. No. 9.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                          13:
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                                                                                                                                                                                   Mismatches
                                                                                                                                                                                   0; Indels
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COUNTRY: USA
ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/954,210
FILING DATE: 20-OCT-1997
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: McMastere, David D.
REGISTRATION NUMBER: 3,963
REFERENCE/DOCKET NUMBER: 3,963
REFERENCE/DOCKET NUMBER: 480124.403C1
TELEPAN: (206) 622-4900
TELEFAX: (206) 622-691
INFORMATION FOR SEQ ID NO: 11:
SPONENCE CHARACTERISTICS:
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           COUNTRY: USA
ZIP: 94022
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/618,834C
FILING DATE: 19-MAR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 5, Application US/08618834C Patent No. 6361937 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 3.8%; Score 12; DB 3; Best Local Similarity 100.0%; Pred. No. 9.8e+03 Matches 12; Conservative 0; Mismatches 0
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                                                                                                                                                                                                                                                                         APPLICANT: Stryer, Lubert
TITLE OF INVENTION: Computer-Aided Nucleic
TITLE OF INVENTION: Sequencing
NUMBER OF SEQUENCES: 54
CORRESPONDENCE ADDRESS:
ADDRESSEE: Ritter, Van Pelt & Yi LLP
STREET: 4906 El Camino Real, Suite 205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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APPLICANT: Yu, Mang
TITLE OF INVENTION: HEPATITIS C VIRUS
NUMBER OF SEQUENCES: 73
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CITY: Seattle
STATE: Washingt
                                                                                                                                                                                                                                                         STREET: 4906 El
CITY: Los Altos
                                                                                                                                                                                                                                       STATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          E: SEED and BERRY LLP
6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RIBOZYMES
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RESULT 33
US-09-371-772B-5650/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TYPE: DNA
; ORGANISM: Hepatitie C Virus
US-09-431-419A-11
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Sequence 5650, Application US/09371772B

Patent No. 6566127

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Pavco, Pam

APPLICANT: McSwiggen, Jim

APPLICANT: Etinchcomb, Dan

APPLICANT: Escobedo, Jaime

TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions |

TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor

FILE REFERENCE: MBHB00, 876-U (3237/198)

CURRENT APPLICATION NUMBER: US/99/371,772B

CURRENT FILING DATE: 1999-08-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 11
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Best Local Similarity
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Best Local Similarity 100.0%; Pred. No. 9.8e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Barber, Jack R.
APPLICANT: Welch, Peter J.
APPLICANT: Tritz, Richard
APPLICANT: Yei, Soonpin
APPLICANT: Yu, Mang
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APPLICANT: Yu, Mang
TITLE OF INVENTION: HEPATITIS C VIRUS RIBOZYMES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: 480124.403C3
CURRENT APPLICATION NUMBER: US/09/431,419A
CURRENT FILING DATE: 1999-11-01
NUMBER OF SEQ ID NOS: 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 650-903-3501 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 16
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REFERENCE/DOCKET NUMBER: AF
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-903-3500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION: NAME: Ritter, Michael J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                  12;
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nucleic acid
DEDNESS: single
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5. 9.8e+03;
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RESULT 34
US-08-704-473-8
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                                                                                                                                                                                                                                                                                                       TELEPHONE: 212-790-9090
TELEPAX: 212-869-8864
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYDE. THE STATE OF THE STATE OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 8,
Patent No. 1
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   Matches
                                    Best Local
                                                                 Query Match
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APPLICANT: Gelb, Bruce
APPLICANT: Chapman, Harold
APPLICANT: Desnick, Robert
APPLICANT: Terrory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
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TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
ZIP: 10036/2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: Coruzzi, Laura A.
REGISTRATION NUMBER: 30,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-790-9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: FastSEQ Version 2.0 CURRENT APPLICATION DATA:
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                                                                                                                                                                                  MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR TITLE OF INVENTION: THE TREATMENT OF BONE RESORP TITLE OF INVENTION: INCLUDING OSTEOPOROSIS
                                                                                                                                                                                                            STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/704,473
FILING DATE: 28-AUG-1996
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM:
SOFTWARE: FastSE(
                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET:
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Similarity 100.0%; Pred. No.
12; Conservative 0; Mismatc
                                        Similarity
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1155 Avenue of the Americas
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IBM Compatible
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                                3.8%; Score 12;
100.0%; Pred. No.
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   Mismatches
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                                                                     DB 2;
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                                        9.9e+03
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nes 0;
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                                                                 Length 17;
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                                               Sequence 4, Application US/08909742
Patent No. 6007991
GENERAL INFORMATION:
APPLICANT: Vimala S. Sivaraman
APPLICANT: Hsien-Yu Wang
                                                                                                                                                                                                                                                                                                         Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (516) 822-3582
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Heien-Yu Wang
APPLICANT: Craig C. Malbor
TITLE OF INVENTION: ACTIVI
TITLE OF INVENTION: BREAS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 08/831,994
FILING DATE: April 1, 1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/827,520
FILING DATE: March 28, 1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Word Perfect 6.1 for windows
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: SIRETCHO CITY: Jericho CITY: New York
 APPLICANT: Craig C. TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                           ANTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                                        HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Adams, Lindsay S.
REGISTRATION NUMBER: 36,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/909,742 FILING DATE: August 12, 1997
                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
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b. 6007991
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12; Conservative
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350 Jericho Turnpike
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                                                                                                                                                                                                                                                                                                                                                                                                          linear
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Malbon
ANTISENSE OLIGONUCLEOTIDES FOR MITOGEN-
ACTIVATED PROTEIN KINASES AS THERAPY FOR
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                                                                                                                                                                                                                                                                                                          3.8%;
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SER: 178-225 CIP II
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Pred. No.
                                                                                                                                                                                                                                                                                                         DB 3;
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                                                                                                                                                                                                                                                                                                     Patent No.
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 514
PRIOR APPLICATION DATA:
RAPLICATION UNMER: 08/831,994
FILING DATE: April 1, 1997
PRIOR APPLICATION UNMER: 08/827,520
APPLICATION NUMBER: 08/827,520
FILING DATE: MATCH 28, 1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                APPLICANT: Barber, Jack R
APPLICANT: Welch, Peter J
APPLICANT: Tritz, Richard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Word Perfect 6.1 for windows
                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP
STREET: 6300 Columbia Center,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 17
TELECOMMUNICATION INFORMATION:
TELEPHONE: (516) 822-3550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                 APPLICANT: Barber,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HYPOTHETICAL:
ANTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                  ADUM: 500
STREET: 500
STRY: Seattle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE:
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                                                                                                                                                                  UMBER OF SEQUENCES:
                                                                 STATE: W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Adams, Lindsay S. REGISTRATION NUMBER: 36,425
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: FILING DATE: August
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
ZIP: 11753
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                                                                                                                                                                              LICANT: Yei, Soonpin
LICANT: Yu, Mang
LE OF INVENTION: HEPATITIS C VIRUS RIBOZYMES
                                                                                                                                                                                                                                                                                         3 71, A5
No. 6043077
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                                                   98104-7092
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                                                                                Washington
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                                                                    USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   August 12, 1997
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100.0%; Pred. No.
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                                                                                                                    701 Fifth Avenue
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APPLICANT: JARVIS, THALE
APPLICANT: MCSWIGGEN, JAMES A.
APPLICANT: STINCHCOMB, DAN T.
TITLE OF INVENTION: ENZYMATIC NUCLEIC ACID TREATMENT OF DISEASES
ITITLE OF INVENTION: OR CONDITIONS RELATED TO LEVELS OF C-FOS
FILE REFERENCE: 231/175
CURRENT APPLICATION NUMBER: US/08/998,099A
CURRENT FILING DATE: 1997-12-24
EARLIER APPLICATION NUMBER: 60/037,658
EARLIER APPLICATION NUMBER: 08/373,124
EARLIER FILING DATE: 1997-01-3
EARLIER FILING DATE: 1995-01-13
EARLIER FILING DATE: 1994-05-18
NUMBER OF SEQ ID NOS: 375
SOFTWALE: FREESEQ for Windows Version 3.0
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                                                                                                    RESULT 39
US-08-998-099-134
                                                                                                                                                                                                                                                                                                                                                  ; TYPE: RNA
; ORGANISM: Homo sapiens
US-08-998-099-133
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INFORMATION FOR SEQ ID NO: 71:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                     Query Match
Best Local Similarity 75.0
Conservative
                     Sequence 134, Application US/08998099A Patent No. 6103890 GENERAL INFORMATION:
APPLICANT: JARVIS, THALE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 133, Appl
Patent No. 6103890
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FILING DATE: 20-OCT-1997
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: MCMASTERS, DAVID D.
REGISTRATION NUMBER: 33,963
REFERENCE/DOCKET NUMBER: 480124.403C1
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
APPLICANT: JARVIS, THALE APPLICANT: MCSWIGGEN, JAMES A.
                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 17
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TELEPHONE: (206)
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                                                                                                                                                                                                                            155 TCCTTCCCCAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             141 GAGGTTGCGACC 152
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n Release #1.0,
                                                                                                                                                                                                                                                                                   Score 12; [
Pred. No. 9.
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o. 9.9e+03;
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                                                          RESULT 41
US-09-412-289-4
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CURRENT APPLICATION NUMBER: US/08/998,099A
CURRENT FILING DATE: 1997-12-24
EARLIER APPLICATION NUMBER: 60/037,658
EARLIER FILING DATE: 1997-01-23
EARLIER FILING DATE: 1997-01-23
EARLIER FILING DATE: 1997-01-13
EARLIER FILING DATE: 1995-01-13
EARLIER APPLICATION NUMBER: 08/245,466
EARLIER APPLICATION NUMBER: 08/245,466
EARLIER FILING DATE: 1994-05-18
NUMBER OF SEQ ID NOS: 375
SOFTWARE: FREESEQ for Windows Version 3.0
SEQ ID NO 134
                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Sivaraman, Vimala S.
APPLICANT: Wang, Hsien-Yu
APPLICANT: Wang, Hsien-Yu
APPLICANT: Mang, Hsien-Yu
APPLICANT: Malbon, Craig C.
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDES FOR MITOGEN-ACTIVATED
TITLE OF INVENTION: PROTEIN KINASES AS THERAPY FOR CANCER
FILE REFERENCE: Seq. 1-4 (178-225 CIP II/CON)
CURRENT APPLICATION NUMBER: US/99/412,289
CURRENT FILING DATE: 1999-10-05
CURRENT FILING DATE: 1999-10-05
EARLIER FILING DATE: 1997-08-12
EARLIER FILING DATE: 1997-08-12
EARLIER APPLICATION NUMBER: 08/831,994
EARLIER APPLICATION NUMBER: 08/827,520
EARLIER APPLICATION NUMBER: 08/827,520
EARLIER FILING DATE: 1997-03-28
NUMBER OF SEQ ID NOS: 4
NUMBER OF SEQ ID NOS: 4
NUMBER OF SEQ ID NOS: 4
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; TYPE: RNA
; ORGANISM: Homo sapiens
US-08-998-099-134
                                                                                                                                                                                     US-09-412-289-3
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US-09-412-289-3
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Best Local Similarity 75.0%;
Matches 9; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                          Matches
                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: STINCHCOMB, DAN T.
TITLE OF: INVENTION: ENZYMATIC NUCLEIC ACID TREATMENT OF DISEASES
TITLE OF: INVENTION: OR CONDITIONS RELATED TO LEVELS OF C-FOS
FILE REFERENCE: 231/175
                                                                                                                                                                                                                                                                                                    LENGTH: 17
TYPE: RNA
                                                                                                                                                                                               ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: synthesized
OTHER INFORMATION: antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          155 TCCTTCCCCAGC 166
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5. 6271210
                                         39 cecceccecces 50
                                                                                      12;
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                                                                                        Conservative
                                                                                                              3.8%; Score 12; DB 3; I
100.0%; Pred. No. 9.9e+03;
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Pred. No. 9.9e+03;
                                                                                        Mismatches
                                                                                        0,
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GENERAL INFORMATION:

Sequence 4, Application US/09412289 Patent No. 6271210

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RESULT 43
US-09-343-698-1/c
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                                                                                                                                                                                                                                                                                                                                               US-09-431-419A-71
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US-09-431-419A-71/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDES FOR TITLE OF INVENTION: PROTEIN KINASES AS THERAPY FOR FILE REFERENCE: Seq. 1-4 (178-25 CIP II/CON) CURRENT APPLICATION NUMBER: US/99/412,289
CURRENT FILING DATE: 1999-10-05
EARLIER APPLICATION NUMBER: 08/909,742
EARLIER FILING DATE: 1997-08-12
EARLIER FILING DATE: 1997-08-12
EARLIER APPLICATION NUMBER: 08/831,994
EARLIER APPLICATION NUMBER: 08/827,520
EARLIER APPLICATION NUMBER: 08/827,520
EARLIER FILING DATE: 1997-03-28
NUMBER: DATE: 1997-03-28
NUMBER: DATE: 1997-03-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-412-289-4
                                       Sequence 1, Application US/09343698 Patent No. 6475486 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 71
LENGTH: 17
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GENERAL INFORMATION:
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Barber, Jack R.
APPLICANT: Welch, Peter J.
APPLICANT: Tritz, Richard
APPLICANT: Yei, Soonpin
APPLICANT: Yu, Mang
TITLE OF INVENTION: HEPATITIS C VIRUS RIBOZYMES
FILE REFERENCE: 480124.403C3
                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/431,419A
CURRENT FILING DATE: 1999-11-01
NUMBER OF SEQ ID NOS: 73
                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                       TYPE: RNA
ORGANISM: Hepatitis C Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Wang, Hsien-Yu
APPLICANT: Malbon, Craig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: synthesized
OTHER INFORMATION: antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Sivaraman, Vimala S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
                                                                                                                                                                                                                                                                                  Local
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Local Similarity 100.0%; Pred. No. 9.9e+03;
APPLICANT: Seeman, Gerhard
Bosslet, Klaus
                                                                                                                                                                                                                        141 GAGGTTGCGACC 152
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100.0%; Pred. No. 9.9e+03;
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US-09-343-698-2
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                                                                                                                                                                                                                                                                                                                    Sequence 2, Application US/09343698 Patent No. 6475486 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/09/343,698
FILING DATE: 30-Jun-1999
CLASSIFICATION: <URKnown>
PRIOR APPLICATION DATA:
'APPLICATION NUMBER: 08/325,955
FILING DATE: <UNKnown>
ATTORNEY/AGENT INFORMATION:
NAME: Ogden, Stasia L.
REGISTRATION NUMBER: 36,228
REFERENCE/DOCKET NUMBER: 05552.0981-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
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Hoffman, Dieter
Sedlacek, Hans-Harald
Sedlacek, Hans-Harald
TITLE OF INVENTION: Glycosyl-Etoposide Prodrugs, A Process For
Preparation Thereof And The Use Thereof in Combination '
Functionalized Tumor-Specific Enzyme Conjugates
                                                                                                                   Hoffman, Dieter
Sedlacek, Hans-Harald
TITLE OF INVENTION: Glycosyl-Ecoposide Prodrugs, A Process For
Preparation Thereof And The Use Thereof In Combination With
Functionalized Tumor-Specific Enzyme Conjugates
                                            NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-408-4000
TELEFAX: 202-408-4400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 20005

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                  APPLICANT: Seeman, Gerhard
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                 Dunner
STREET: 1300 I Street,
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TYPE: nucleic acid
STRANDEDNESS: single
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 1300 I Street, N.W., Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                  Czech, Joerg
Kolar, Cenek
                                                                                                                                                                                                                                                                              Bosslet, Klaus
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100.0%; Pred. No. 9.9e+03;
tive 0; Mismatches 0;
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RESULT 45
US-09-371-772B-4397
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                                                                                                                                                                                                                                     ; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-371-772B-4397
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INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn version 3.0 SEQ ID NO 4397
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 4397, Application US/09371772B Patent No. 6566127
GENERAL INFORMATION:
                                                                         Query Match
Best Local Similarity
Matches 10; Conserva
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00,876-J (237/198)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR PELICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Pavco, Pam APPLICANT: McSwlggen, Jim
                                                                                                                                                                                                                                                                                                                                                             LENGTH: 17
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC COMPUTER:

COMPUT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 08/325,955
FILING DATE: <Unknown>
ATTORNBY/AGENT INFORMATION:
NAME: Ogden, Stasia L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Ogden, Stasia L.
REGISTRATION NUMBER: 36,228
REFERENCE/DOCKET NUMBER: 05:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA
    119 GAGTGGACCATC 130
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                                                                              Conservative
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                                                                                                              3.8%;
                                                                                                                  Score 12; DB 4;
Pred. No. 9.9e+03;
                                                                                  Mismatches
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RESULT 47
US-08-325-955-2
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; MOLECULE TYPE:
US-08-325-955-1
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                                                       GENERAL INFORMATION:
APPLICANT: Seeman, Gerhar
APPLICANT: Bosslet, Klaus
APPLICANT: Czech, Joerg
APPLICANT: Kolar, Cenek
APPLICANT: Hoffman, Diete
APPLICANT: Sedlacek, Hans
                                                                                                                                                                                         Sequence 2, Applic
Patent No. 6610299
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                                                                                                                                                                                                                                                                                                                                                                                              Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: Ogden, Stasia L.
REGISTRION NUMBER: 36,228
REFERENCE/DOCKET NUMBER: 0555
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202.408-4000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/325,955
FILING DATE: 19-0CT-1994
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: G1:
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                APPLICANT: Hoffman, Dieter
APPLICANT: Sedlacek, Hans-Harald
TITLE OF INVENTION: Glycosyl-Etoposide Prodrugs, A Process For
NUMBER OF SEQUENCES: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                              Local
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ZIP: 20005
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Finnegan, Henderson, Farabow, Garrett & ADDRESSEE: Dunner
STREET: 1300 I Street, N.W., Suite 700
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: HOffman, Dieter

F: Sedlacek, Hans-Harald

INVENTION: Glycosyl-Etoposide Prodrugs, A Process

ROUENCES: 8
                                                                                                                                                                                                                Application US/08325955
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Bosslet, Klaus
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Czech, Joerg
Kolar, Cenek
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hes 0;
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ADDRESSEE:

Finnegan, Henderson, Farabow, Garrett &

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CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: DCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
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US-09-866-108A-8273/c
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PM PC COMPUTER: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, VE
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/325,955
                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: AEOMICA-7
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                PPLICANT: SHANNON, Mark
ITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
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REGISTRATION NUMBER: 36,228
REFERENCE/DOCKET NUMBER: 05:
TELECOMMUNICATION INFORMATION:
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CLASSIFICATION: 435
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CITY: Washington
STATE: D.C.
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                                                                                           APPLICATION NUMBER: PCT/US01/00667
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
                                 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION:
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Similarity 100.0%;
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                NUMBER: PCT/US01/00665
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2001-01-30
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; Pred. No.
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                                                                                 ; ORGANISM: Homo sapiens
US-09-866-108A-8274
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CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
PRIOR PILING DATE: 2000-10-04
PRIOR PILING DATE: 2000-09-27
PRIOR PILING DATE: 2000-09-27
PRIOR PILING DATE: 2000-09-27
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR APPLICATION NUMBER: PCT/US01/00664
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; SEQ ID NO 8273
; LENGTH: 17
; TYPE: DNA
                                                                                                                                                 Patent No. 6686
SEQ ID NO 8274
LENGTH: 17
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                                           Query Match
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APPLICANT:
                                                                                                                                                                                                              Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 15755
SOFTWARE: Aeomica Sequence Listing Engine
                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
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SOFTWARE: Acomica Sequence Listing Engine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: PCT/US01/00668 PRIOR FILING DATE: 2001-01-30 PRIOR APPLICATION NUMBER: PCT/US01/00663 PRIOR FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
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    Local Similarity
nes 12; Conserv
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                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: PCT/US01/00665 FILING DATE: 2001-01-30
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    Conservative
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                     100.0%;
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  3.8%; Score 12; DB 4;
100.0%; Pred. No. 9.9e+(
ive 0; Mismatches
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Pred. No.
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                   9.9e+03;
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                                         Length 17;
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US-09-866-108A-8275/c
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; Sequence 8276, Application US/09866108A
; Patent No. 6686188
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PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR FILING DATE: 2000-09-27

PRIOR PILING DATE: 2001-09-27

PRIOR APPLICATION NUMBER: PCT/US01/00667

PRIOR PILLING DATE: 2001-01-30

PRIOR PILLING DATE: 2001-01-30

PRIOR PILLING DATE: 2001-01-30

PRIOR PRILLING DATE: 2001-01-30

PRIOR PELLICATION NUMBER: PCT/US01/00664

PRIOR PELLICATION NUMBER: PCT/US01/00669

PRIOR PILLING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/US01/00665
                                                                                                                      GENERAL INFORMATION:
APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharro
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APPLICANT: CHEN, Wenshang
APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYGSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR PRIOR DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
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APPLICANT: JI, Yongga
APPLICANT: PENN, Shar
APPLICANT: HANZEL, Da
                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Homo sapiens
                                                       APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
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12; Conserv
F: SHANNON, Mark
INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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                                                                      PENN, Sharron G.
HANZEL, David K.
RANK, David R.
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HANZEL, David K.
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                                                CHEN, Wensheng
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100.0%; Pred. No.
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CURRENT FILING DATE: 2001-05-25

PRIOR APPLICATION NUMBER: US 60/207,456

PRIOR FILING DATE: 2000-05-26

PRIOR PELLING DATE: 2000-05-26

PRIOR APPLICATION NUMBER: GB 24263.6

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR APPLICATION NUMBER: PCT/US01/00666

PRIOR PILING DATE: 2000-09-27

PRIOR APPLICATION NUMBER: PCT/US01/00666

PRIOR PILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-01-30
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; ORGANISM: Homo sapiens
US-09-866-108A-8276
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SEQ ID NO 8276
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APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEDMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: AEOMICA-7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: PCT/US01/00663 PRIOR FILING DATE: 2001-01-30
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                                                     FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
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FILING DATE: 2000-09-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
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                                     APPLICATION NUMBER:
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N, Sharron
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PCT/US01/00665
                                   PCT/US01/00669
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Pred. No.
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9.9e+03;
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FILING DATE: 2001-01-30

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US-09-866-108A-8278/c
                                                                                         ; ORGANISM: Homo sapiens
US-09-866-108A-8278
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SEQ ID NO 8277
          Best Loc
Matches
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ORGANISM: Homo sapiens
-09-866-108A-8277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
                                                                                                                                                                                                               SOFTWARE:
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
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PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ENERAL INFORMATION:
APPLICANT: GU, Yizhong
APPLICANT: JI, Yongga
                                                                                                                                                                                                                             Remaining Prior Application data removed - NUMBER OF SEQ ID NOS: 15755
                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
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PRIOR FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: RANK, David R.
APPLICANT: CHEN, Mensheng
APPLICANT: SHANNON, MAIK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
                                                                                                                                                                                                                                                                           PRIOR FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: PCT/US01/00663
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                                                                                                                                 TYPE: DNA
          y Match 3.8%; Score 12; DB 4; Le Local Similarity 100.0%; Pred. No. 9.9e+03; hes 12; Conservative 0; Mismatches 0;
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                            RESULT 55
US-08-712-357-9/c
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Sequence 9, Application US/08712357 Patent No. 5808037
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                                                                                                                                                                                                                                                                                                                                           TELEFAX: 816-474-9057
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                      TOPOLOGY: linear MOLECULE TYPE: DNA HYPOTHETICAL: NO ANTI-SENSE: YES POSITION IN GENOME: UNITS: bp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                       NAME: Collins, John M.
REGISTRATION NUMBER: 26,262
REFERENCE/DOCKET NUMBER: 24:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 816-474-9050
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/802;547
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET:
                                                                                                                                                                                                                                                                                                                                            TELEPHONE: 816-474-9057
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
CLASSIFICATION:
                                                                                                                                                    Local
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5. 5780611
                                                                                                                                      al Similarity
12; Conserv
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2405 Grand Boulevard, Suite 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            USA
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                                                                                                                                       Conservative
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Kovacs, Attilla
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OLIGOMERS WHICH
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Pred. No.
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                                                                                                                                                                  Length 18;
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                                                                                                                                       Gaps
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GENERAL INFORMATION:
APPLICANT: Guntaka
APPLICANT: Weber,
APPLICANT: Kovacs,

Guntaka, Ramareddy V. Weber, Karl T.

APPLICANT: Kandala, Jagannadhachari TITLE OF INVENTION: OLIGOMERS WHICH INHIBIT

Kovacs, Attila

The Water

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Query Match
Best Local Similarity
Watches 12; Conserve
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                                                                                                                           ; OTHER INFORMATION: Antisense Oligonucleotide US-09-205-204-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-08-712-357-9
                                                                                                                                                                                                         Sequence 40, Application US/09205204
Patent No. 5958772
GENERAL INFORMATION:
APPLICANT: C. Frank Bennett
APPLICANT: Elizabeth J. Ackermann
APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF CELLULAR INHIBITOR OF APOPTOSIS-1 EXPRESS
FILE REFERENCE: RTS-0020
CURRENT APPLICATION NUMBER: US/09/205,204
CURRENT FILING DATE: 1998-12-03
NUMBER OF SEQ ID NOS: 47
SEQ ID NO 40
LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
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TELEFAX: (816) 474-9057
INFORMATION FOR SEQ ID NO:
                                                                                                                                                              FEATURE:
                                                                                                                                                                         ORGANISM: Artificial Sequence
                                                                                                                                                                                                TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ANTI-SENSE: YES
POSITION IN GENOME:
UNITS: bp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: COLLINS, John M.
REGISTRATION NUMBER: 26262
TELECOMMUNICATION INFORMATION:
TELEPHONE: (816) 474-9050
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/712,357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HYPOTHETICAL:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 64108
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS:
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   263 AACAAGCTGAAG 274
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RET: 2405 Grand Boulevard, Suite 400
K. Kaneas City
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          : 18 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Missouri
                                                  3.8%; Score 12; DB 2; Lot 12; DB 2; Lot 13; Pred. No. 9.9e+03; Conservative 0; Mismatches 0;
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100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; OTHER INFORMATION: Antisense Oligonucleotide US-09-205-860-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 57
US-09-205-860-10
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APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF G-ALPHA-13 EXPRESSION
FILE REFERENCE: RTS-0031
FILE REFERENCE: RTS-0031
CURRENT APPLICATION NUMBER: US/09/205,860
CURRENT PILING DATE: 1998-12-04
NUMBER OF SEQ ID NOS: 87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Patent No. 5994075
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 10, Application US/09205860 Patent No. 5981732
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Best Local :
                                                                                                     TELEFAX: 617-345-9111
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 18
TYPE: DNA
ORGANIEM: Artificial Sequence
FEATURE:
                                                                                                                                                 REGISTRATION NUMBER: 34,380
REFERENCE/DOCKET NUMBER: 35
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                             CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: LEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
 MOLECULE TYPE:
                                                                                   SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                        APPLICATION NUMBER: US/6
FILING DATE: 17-MAY-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: FILING DATE: 16-MA
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                              STRANDEDNESS:
                                                                                                                                    TELEPHONE: 617-345-9100
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                                                                   ENGTH:
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                                           nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Boston
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                                                                   18 bases
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                    linear
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other nucleic acid
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hes 0;
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APPLICANT: Bennett, C. Frank
APPLICANT: Cowsert, Lex M.
TITLE OF INVENTION: ANTISENSE MODULATION OF CD:
FILE REFERENCE: RTS-0007
CURRENT APPLICATION NUMBER: US/09/161,244
CURRENT FILING DATE: 1998-09-25
NUMBER OF SEQ ID NOS: 91
SEQ ID NO 63
LENGTH: 18
                              RESULT 61
US-08-970-740-14/c
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; TYPE: DNA
; CARNISM: Artificial Sequence
; EEATURE:
; OTHER INFORMATION: Antisense Oligonucleotide
US-09-161-244-63
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US-09-161-244-63/c
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Sequence 14, Application US/08970740 Patent No. 6015670
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GENERAL INFORMATION:
                                                                                                                                                                       Query Match
Best Local
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LENGTH: 18
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Best Local Similarity
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                                                                                                                                                        Matches
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APPLICANT: ex M. Cowsert
TITLE OP INVENTION: ANYLISENSE MODULATION OF EGR-1 EXPRESSION
PILE REFERENCE: RTS-0028
FILE REFERENCE: RTS-0028
FILE REPERENCE: NUMBER: US/09/205,921A
CURRENT FILLING DATE: 1998-12-04
NUMBER OF SEQ ID NOS: 47
                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                OTHER INFORMATION: Antisense Oligonucleotide
                                                                                                                                                                                                                                                                                 TYPE: DNA
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                                                                                                                                                                     3.8%; Score 12; DB 3; 100.0%; Pred. No. 9.9e+(
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Pred. No. 9.9e+03;
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US-09-143-212-44/c
; Sequence 44, Application US/09143212B
; Patent No. 6077672
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                                                                        US-09-143-212-44
Query Match
Best Local Sim
Matches 12;
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APPLICANT: Brett P. Monia and Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF TRADD EXPRESSION
FILE REFERENCE: RTS-0005
                                                                                                                                                             SEQ ID NO 44
LENGTH: 18
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Best Local (
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: WORDPETFECT 6.1
CURRENT APPLICATION UNMBER: US/08/
FILING DATE: 14-NOV-1997
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 08/857
FILING DATE: 16-MAY-1997
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 60/017
APPLICATION UNMBER: 60/017
FILING DATE: 17-MAY-1996
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                                                                                                                         TYPE: DNA ORGANISM: Artificial Sequence
                                                                                        OTHER INFORMATION:
                                                                                                           FEATURE:
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LENGTH: 18 bases
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Goodfellow, P.N.
TITLE OF INVENTION: METHODS FOR IDEN
TITLE OF INVENTION: GENE OF INTEREST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: 1: MOLECULE TYPE:
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CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Kathleen M. Williams REGISTRATION NUMBER: 34,38
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h 3.8%; Score 12; DB Similarity 100.0%; Pred. No. 9.9 12; Conservative 0; Mismatches
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                                                                                      Antisense Oligonucleotide
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100.0%; Pred. No. 9.9
ive 0; Mismatches
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                    DB 3; I
9.9e+03;
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                                   Length 18;
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39

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RESULT 63
US-09-143-212-45/c
; Sequence 45, Application US/09143212B
; Patent No. 6077672
; GENERAL INFORMATION:
; APPLICANT: Brett P. Monia and Lex M.
; TITLE OF INVENTION: nmc.nons
RESULT 65
US-08-679-645-1163
; Sequence 1163, Application US/08679645
; Patent No. 6350934
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 12; Conserve
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                                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: Antisense Oligonucleotide US-09-496-694B-176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-496-694B-176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; OTHER INFORMATION: Antisense Oligonucleotide
US-09-143-212-45
                                                                                                                                                                                                                                Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 45
LENGTH: 18
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 176, Application US/09496694B Patent No. 6335194
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 176
LENGTH: 18
                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF SURVIVIN EXPRESSION
FILE REFERENCE: ISPH-0439
CURRENT APPLICATION NUMBER: US/09/496,694B
CURRENT FILING DATE: 2000-02-02
PRIOR APPLICATION NUMBER: 09/286,407
PRIOR PILING DATE: 1999-04-05
PRIOR PILING DATE: 1998-09-29
PRIOR FILING DATE: 1998-09-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Brett P. Monia and Lex M. Cowsert TITLE OF INVENTION: ANTISENSE MODULATION OF TRADD EXPRESSION FILE REFERENCE: RTS-0005 CURRENT APPLICATION NUMBER: US/09/143,212B CURRENT FILING DATE: 1998-08-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: C. Frank Bennett
APPLICANT: Elizabeth J. Ackermann
APPLICANT: Eric E. Swayze
APPLICANT: Lex M. Cowsert
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                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                      Local
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12; Conser
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Skokut, Thomas A.

Guo,

Lining

McSwiggen, James A. Merlo, Patricia Ann

Owens

US-08-679-645-1165

Application US/08679645

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RESULT 66
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                                                                                                         Matches
                                                                                                                      Best Local
                                                                                                                                   Query Match
                                                                                                                                                                                                                                             TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Zwick, Michael G
APPLICANT: SEdington, Brent
APPLICANT: McSwiggen, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                        NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 21:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/001,135
APPLICATION NUMBER: 08/001,135
APPLICATION NUMBER: 08/300,726
FILLING DATE: September 2, 1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM Compatib.
OPERATING SYSTEM: IBM |
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                          TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/679,645 FILING DATE: July 12, 1996 CLASSIFICATION: 800
                                                                                                                                                                                   STRANDEDNESS:
TOPOLOGY: lin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Los Angeles
                                                                                                                                                                                                              TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
                                                                                                                                                                                                                              ENGTH:
                                                                         39 ceccecceccec so
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                                                                                                                      Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E: Lyon & Lyon
633 West Fifth Street
                                                                                                                                                                                                                              18 base pairs
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                                                                                                        Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SYSTEM: IBM P.C.
Word Perfect 5.1
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                                                                                                                                                                                   linear
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                                                                                                    3.8%; Score 12; DB 3; L
100.0%; Pred. No. 9.9e+03;
Live 0; Mismatches 0;
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                                                                                                                                   Length 18;
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Young, Scott A.

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                                                                                                                                                                                                                                                                      Sequence 21, Application US/09063733A Patent No. 6372211
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                               GENERAL INFORMATION:
APPLICANT: Isaac, Barbara G.
APPLICANT: Iseac, John T.
APPLICANT: Purcell, John P.
APPLICANT: Purcell, John P.
APPLICANT: Romano, Charles P.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR CONTROLLING
TITLE OF INVENTION: INSECTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/679,645
FILING DATE: July 12, 1996
CLASSIFICATION: 800
PRIOR APPLICATION NUMBER: 60/001,135
FILING DATE: July 13, 1995
APPLICATION NUMBER: 08/300,726
FILING DATE: July 13, 1995
APPLICATION UNMBER: 08/300,726
FILING DATE: September 2, 1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Merlo, Donald J.
TITLE OF INVENTION: COMPOSI-
TITLE OF INVENTION: MODULAN
TITLE OF INVENTION: IN PLAN
NUMBER OF SEQUENCES: 1263
                                                                                        NUMBER OF SEQUENCES: CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
COUNTRY:
                      STATE:
                                                      STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 219/247
                                                                        ADDRESSEE:
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12; Conservative (
                Houston
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                                                        PO Box 4433
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                                                                        Arnold White & Durkee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3.8%;
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US-09-679-298A-25
US-09-679-298A-25
Sequence 25, Applicati
Patent No. 6566131
GENERAL INFORMATION:
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                                                                                                                                            RESULT 69
US-09-077-940A-8/c
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GENERAL INFORMATION:
APPLICANT: KIMURA, TOTH et al.
TITLE OF INVENTION: NOVEL SEMAPHORIN Z AND
FILE REFERENCE: 0020-4426P
CURRENT APPLICATION NUMBER: US/09/077,940A
CURRENT FILING DATE: 1998-06-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 47
SEQ ID NO 25
LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 713-787-1440
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                      Matches
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                                                                                                          Sequence 8, Application US/09077940A Patent No. 6576441
                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Brett P. Monia
APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF SMAD6 EXPRESSION
FILE REFERENCE: RTS-0045
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/679,298A CURRENT FILING DATE: 2001-03-05
                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Antisense Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Patterson, Melinda L. REGISTRATION NUMBER: 33,062 REFERENCE/DOCKET NUMBER: MO TELECOMMUNICATION INFORMATION: TELEPHONE: 713-787-1400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: 715-,.
TELEPHONE: 713-787-1440
TO TO NO:
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                                                                                                                                                                                                                                                                                                         Local Similarity
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Pred. No.
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                                                                                                                                                    RESULT 71
US-08-299-187-6
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                                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: (GCG)6 probe US-09-500-700-68
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 08/676,318
PRIOR FILING DATE: 1996-12-30
PRIOR APPLICATION NUMBER: PCT/US95/00829
PRIOR FILING DATE: 1995-01-18
PRIOR APPLICATION NUMBER: US 08/312,604
PRIOR FILING DATE: 1994-09-28
PRIOR PFLICATION NUMBER: US 08/183,119
PRIOR FILING DATE: 1994-01-18
                                                                                                                sequence 6, Application US/08299187
Patent No. 5736325
                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn version 3.1 SEQ ID NO 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 12; Conserv
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 68, Application US/09500700 Patent No. 6790941
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                                                                                                                                                                                                                                                                                           Matches
                                                               GENERAL INFORMATION:
APPLICANT: Manowitz, Paul
APPLICANT: Poretz, Ronald D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT FILING DATE: 2003-01-10 PRIOR APPLICATION NUMBER: US 08/863,813 PRIOR FILING DATE: 1997-05-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: SCRIP1160-4
CURRENT APPLICATION NUMBER: US/09/500,700
CURRENT FILING DATE: 2003-01-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: THE SCRIPPS RESEARCH INSTITUTE APPLICANT: BARBAS III, Carlos F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: WRIGHT, Peter E.
TITLE OF INVENTION: ZINC FINGER PROTEIN DERIVATIVES AND METHODS THEREFOR
                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                              ENGTH:
                                   APPLICANT:
   ITLE OF INVENTION:
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                                                                                                                                                                                                                                                        39 ceccecceccec so
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                                 Park, David
Ricketts, Michael H.
                                                                                                                                                                                                                                                                                          Conservative
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100.0%; Pr/
MARKER FOR INDIVIDUALS SUSCEPTIBLE TO ALCOHOLISM
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100.0%; Pred. No.
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0; Mismatches
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                                                                                                                                                                                                                                                                                                         9.9e+03
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                                                                                                                                                                                          Sequence 16, Application US/08715142 Patent No. 5811244
                                                                                                                                                                                                                                                                                                                                                    Matches
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TELEX: 133521
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                APPLICANT: France,
APPLICANT: Cox, Gregory A.
APPLICANT: Lutz, Cathleen M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORIGINAL SOURCE:
ORGANISM: Homo sapiens
PUBLICATION INFORMATION;
AUTHORS: Gieselmann, Volkmar
                                                           APPLICANT: Lutz, Cathleen M.
APPLICANT: No. 5811244bols, Jeffrey L.
TITLE OF INVENTION: CLINICAL DISORDERS
TITLE OF INVENTION: MUTATION
NUMBER OF SEQUENCES: 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: 201 487-5800
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CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                              CORRESPONDENCE ADDRESS:
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DATE: December-1989
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Jackson Esq., David A.
REGISTRATION NUMBER: 26,742
REFERENCE/DOCKET NUMBER: 601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AUTHORS: von Figura, Kurt TITLE: Arylaulfatase A pseudodeficiency: Loss of TITLE: polyadenylation signal and N-glycosylation JOURNAL: Proc. Natl. Acad. Sci. U.S.A.
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TOPOLOGY: lin
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                               ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                     222 AGGTGACACTGG 233
                                                                                                                                                                                                                                                                                                                                                    12;
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York Harbor
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                E: Kevin M.
P.O. Box 999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
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                               Farrell,
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                                                                                              ASSOCIATED
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COUNTRY: U

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                                                                                               COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/88/465,485A
FILING DATE: 05-JUN-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/124,256
PILING DATE: 20-SEP-1993
PRIOR APPLICATION NUMBER: US 07/840,716
APPLICATION NUMBER: US 07/840,716
FILING DATE: 21-FEB-1992
PRIOR APPLICATION NUMBER: US 07/840,716
FILING DATE: 21-FEB-1992
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GENERAL INFORMATION:
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TELEFAX: (207) 363-0528
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 1/50
CITY: Arlington
STATE: Virginia
STATE: U.S.A.
                   APPLICATION NUMBER: US 07/840,716
FILING DATE: 21-FEB-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/288,692
FILING DATE: 22-DEC-1988
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER REALABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: PatentIn Release #1.0, Version #1.25
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/08/715,142
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LENGTH: 20 base pairs
TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
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REGISTRATION NUMBER: 35,505
REFERENCE/DOCKET NUMBER: JL-9601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/0 FILING DATE: 18-SEP-1996 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: P.C.
STREET: 1755 S. Jefferson Davis Hwy., Suite 400
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Fortney, Andrew D
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100.0%; Pred. No. 9.9e+
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Best Local
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INFORMATION FOR SEQ ID NO: 28:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                      TELEX: 133521
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 60
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-487-5800
                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/910,443
FILING DATE: 05-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (408) 436-2070
                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: Jackson Esq., David A.
REGISTRATION NUMBER: 26,742
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
HYPOTHETICAL:
           MOLECULE TYPE: other nucleic acid DESCRIPTION: /desc = "PRIMER"
                                                                                                       SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
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                                                        TYPE: nucleic acid
STRANDEDNESS: sing
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100.0%; Pred. No.
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Sequence 11, Application US/09358382

Patent No. 6010906

GENERAL INFORMATION:
APPLICANT: Donna T. Ward
APPLICANT: Lex M. Cowsert
FILE REFERENCE: RTS-0071

CURRENT PELLICION UNMEER: US/09/358,382

CURRENT FILING DATE: 1999-07-21

NUMBER OF SEQ ID NOS: 47

FYDE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Antisense Oligonucleotide

US-09-358-382-11
Search completed: February 2, 2005, 23:37:48 Job time: 33.5987 secs
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US-09-358-382-11
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AZ603262 1M0422P24 AZ833326 ZM0115915 CF298174 7LEAF-01 AJ651227 AZ503426 1M0003A09 AZ863893 ZM0180117 AZ477495 1M0209A06 AL491155 T. brucei AZ759923 1M0553110 AZ867011 ZM0103112 CL438501 PST7730-N AZ389514 1M0150108 AZ610580 1M045N21 AL481612 T. brucei AA916934 cn14409.s AZ820818 ZM0093118 AJ666323 AJ666323 AZ780308 ZM0017J05 AZ333915 1M0063002 AZ431118 1M019D16 A1035419 Ub46405.r AZ810074 ZM0074J19 AZ586137 1M0391M17 AZ829971 ZM0107K20	by chance to have a he result being printed, distribution. Description		46458		ctacctgccgctgtcccgtg 132	en Ltd. h time 405.351 Seconds allgnments) Million cell updates/sec
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AZ603262
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   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleo Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murin 1 (bases 1 to 20)
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T.,
                                                                                                                              AZ603262 20 bp DNA linear GSS 13-DEC-200
1M0422P24F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0422P24 F, genomic survey sequence.
                                                                    Mus musculus
                                                                               Mus musculus (house mouse)
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AI040238
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AII4974493
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                                            Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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R54679 yj74e12.r1
W92724 zd92a07.s1
AZ307173 1M0008N03
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AZ486627 1M0314H13
AZ579398 1M0359B2
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H84968 ys89c08.r1
AZ442177 1M0234K14
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AJ668095
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AL483711
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                   Hamil, C.,
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T. brucei

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                      AUTHORS
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                                                                                                                                                                                                                                                                                   Local
Mammalia, Eutheria; Rodentia; C:
Mammalia, Eutheria; Rodentia; Sc
1 (bases 1 to 27)
Dunn,D., Aoyaqi,A., Barber,M., B.
Islam,H., Longacre,S. Mah-
                                                                                                                                                                                                                                    104 CCCTGTGTCCTA 115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.e
Insert Length: 10000 Std E
Plate: 0422 row: p column:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
High onalism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           inpublished (2000)
Contact: Robert B. Weiss
University of Utah
University of Utah
Rm. 308_Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Niederhausern, A. ar
Mouse whole genome
plasmid inserts
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12; Conserv
                                                                                                                      AZ833326 27 bp DNA linear GSS 20-FEB-200 2M0115915F Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGC2M0115P15 F, genomic survey sequence.
AZ833326
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                                                                                 musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwD42 (gi[4732114]gb[AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                              (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"/Clone lib="Mouse 10kb plasmid UUGCIM library"/note="Vector: pW042nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Laboratory Mouse DNA Resource
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/strain="C57BL/6J"
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e genome scaffolding with paired end reac
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       rber,M., Beacorn,T., Duv
Mahmoud,M., Meenen,E.,
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Duval,B.,

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Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Cyza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.

1 (bases 1 to 25)
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                                                                                                                                                                                                                               7LEAF--01-H06.gl sativa (japonica
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                                                                                                                                                                                                                                                                                CF298174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Err
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Fax: 801 585 7177
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Contact: Robert B.
                                                                                                                                                                 CF298174.1
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi 4732114 |gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /lab_host="B. Coli strain XL10-Gold, T1-resistant, F-"/Clone_lib="Mouse 10kb plasmid UUGCIM library"/note="Vector: pW042nv; Purified genomic DNA from M.musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Laboratory Mouse DNA Resource
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/strain="C57BL/6J"
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Rice leaf plasmid cDNA library II (7LEAF) Oryza
cultivar-group) cDNA clone 7LEAF--01-H06, mRNA
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
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Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae; Sus.
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Similarity 100.0%; Pred. No. 7.7e+05;
11; Conservative 0; Mismatches 0;
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/dev_stage="7 days after germination"
/dev_stage="7 days after germination"
/lab_nost="E.coli DH10B"
/clone_lbb="Rice_leaf_plasmid_cDNA_library_II_(7LEAF)"
/clone_lbb="Rice_leaf_plasmid_cDNA_library_II_(7LEAF)"
/note="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA_was_capped
/mth_oligoribonucleotides_and_then_used_as_templates_for_mrna_ref_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_prince_leaf_
/tissue_type="ovary"
/clone_lib="CSEQRAN19"
/note="Vector: pBlueScript11(KS+); Site_1: EcoRI; Site_2:
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                                                                                                                                                                                          /clone="C0003277_L02"
                                                                                                                                                                                                                                          mol_type="mRNA"
db_xref="taxon:9823"
                                                                                                                                                                                                                                                                                                                                                                   organism="Sus scrofa"
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/cultivar="Nackdong"
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clone="7LEAF--01-H06"
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AZ303426
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Seq primer: CGTTGTAAAACGACGGCCAGTClass: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0003 row: A column: 09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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AZ303426.1 GI:10338804
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (2000)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mus musculus
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801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         USA
                                                                               10.5 kb range using preparative agarose gel lectrophoresis. Vector DNA was prepared from a derivative of pND42 (gi|473214|gb|AFT29072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated
                                                                                                                                                                                        (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                                                                                                                                                                                                 musculus C57BL/6J (male) was obtained Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Biomedical
                                                        with adaptors complementary to the insert adaptors and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NotI; Single pass sequencing; Normalised library constructed from pooled ovaries"
                                                                                                                                                                                                                                                                                                                                                                                                                                       /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            'mol_type="genomic DNA"
'strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sex="Male
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         clone="UUGC1M0003A09"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               db_xref="taxon:10090"
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The sheared, adaptored mouse DNA was annealed to vector DNA, and transformed into
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Polymers Research Bldg., 20
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AZ868893
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0180 row: I column: 17
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    University of Utah University of Utah
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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High quality sequence stop: 27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished (2000)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 308,
(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           USA
                                                                                                                                                                                                                                                                                                                                   /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/Clone lib="Mouse 10kb plasmid UUGCIM library"/note="Vector: PWD42TV; Purified genomic DNA from M.musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Biomedical Polymers Research Bldg., 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                       clone="UUGC2M0180I17"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                organism="Mus musculus"
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                                                                                                                                                                                                                                                                                                                                                                                                                                   sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Fr
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0; Mismatches
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3. 7.7e+05;
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AZ427495
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Seq primer: CACACAGGAAACAGCTATGACC Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0209 row: A column: 06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 University of Utah Genome Center University of Utah Rm. 308, Biomedical Polymers Res
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (2000)
Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R.
Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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AZ427495.1 GI:10551508
GSS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         quality sequence stop: 28.
Location/Qualifiers
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              Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 bb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD42 (gi|4732114|gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mol_type="genomic DNA"
strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               clone="UUGC1M0209A06"
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and transformed into
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us genomic
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RESULT 9
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AUTHORS
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TA316B05P/c
                                                            DEFINITION
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JOURNAL
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Best Local
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                                                                                                                                                                                                                             54 CATGCACACGT 64
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Chamber ac.uk

Constructed at the Institute for Genomic Research (TICR),

Constructed at the Institute for Genomic Research

Rockville, MD. Genomic DNA isolated from a cloned population of

Rypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically sheared

to give a tight size distribution (

4 kb). The v + i method used for the library construction is

described in detail in Smith, H. and Venter, J.C. (Making small

insert libraries for whole genome shotgun sequencing projects. In

Genome Sequencing: A Practical Approach, eds. M. Vaudin and B.

Barrell, Oxford University Press, 1999).
                                                                                                                                                                                                                                                                        11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Direct Submission
Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hit Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and
                                       1M0553I10F Mouse 10kb plasmid clone UUGC1M0553I10 F, genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger Centre are available at http://www.sanger.ac.uk/Projects/T_brucei/.
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                                                                                  AZ759923
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Trypanosoma
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                       AZ759923
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                                                                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                                                                                                                                                        /mol_type="genomic DNA"
/strain="TREU927"
/db_xref="taxon:5691"
/clone="316b05"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           organism="Trypanosoma brucei"
GI:12867209
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100.0%; Pred. No. 7.7e+05;
tive 0; Mismatches 0;
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Mus musculus (house mouse)

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KEYWORDS
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Eukaryota; Metazoa; Chordata;
Eutheria; Rodentia;
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11; Conserv
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                                                                                                AZ827011 29 bp DNA linear GSS 20-FEB-200200103112F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC2M0103112 F, genomic survey sequence.
                                                    AZ827011.1 GI:12996919
                                                                                  AZ827011
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Insert Length: 10000 Std Error: 0.00
Plate: 0553 row: I column: 10
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Fax: 801 585 7177
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University of Utah Genome Center
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E.,
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igh quality sequence stop: 29.
Location/Qualifiers
musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNAP2 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/clone lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: pWD42nv purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackso
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/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0553I10"
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                                                                          CL438501 29 bp DNA linear GSS 18-MAR-2004
PST7630-NL.Seq MICB1 Mus musculus genomic clone PST7630-NL.Seq,
genomic survey sequence.
CL438501
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High quality sequence stop: 29.
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Insert Length: 10000 Std Error:
Plate: 0103 row: I column: 12
Seq primer: CGTTGTAAAACGACGGCCAGT
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Contact: Robert B.
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Fax: 801 585 7177
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/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC2M0103I12"
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100.0%; Pred. No. 7.7e+05;
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UNNeoSVI gene trap. Tag generated by plasmid rescue. Additional
sequence information and target gene cloning can be generated. Es
cell line harboring insertion mutation of target gene is available.
Sequence analysis available from
                                                                                                                                                                                                                                                                 Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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1M0150108F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0150108 F, genomic survey sequence.
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Manitoba Institute of Cell Biology, Universitry of Manitoba
CN5029, 675 McDermot Ave, Winnipeg, MB R3E 0V9, Canada
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Unpublished (2002)
                                                Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                     University of Utah Genome Center University of Utah
                                                                                                                                                                                   Contact: Robert B. Weiss
                                                                                                                                                                                                        Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                              Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AZ389514.1 GI:10503222
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Fax: 204 787 2190
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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ilarity 100.0%;
Conservative (
  ddunn@genetics.utah.edu
Length: 10000 Std Erry
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                                                                                                              Biomedical Polymers Research Bldg.,
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/cell_line="D3H (J1 subclone)"
/clone_lib="MICB1"
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/mol_type="genomic DNA"
/strain="129 sv"
/db_xref="taxon:10090"
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R. Mouse whole genome scaffolding with paired end reads from 10kb mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                            University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Res
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1M0435N21R Mouse 10kb plasmid clone UUGC1M0435N21 R, genomic
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Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
                                                                                                                                                                                 Contact: Robert B.
                                                                                                                                                                                                 plasmid inserts
Unpublished (2000)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pNP42 (gi|4732114|gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coli XI10-Gold (Stratagene) cells
and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
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strain="C57BL/6J"
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                                                                                                              Polymers Research
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Std Error: 0.00
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UUGC1M library Mus musculus genomic
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Best Local :
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Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a clomed population of Trypanosoma brucei (TREU927/4 GUTTAT 10.1) was mechanically sheared to give a tight size distribution (4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. In Genome Sequencing: A Practical Approach, eds. M. Vaudin and B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11;
                                                                                                                                                                                     Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hi Cambridge CB10 18A, E-mail: barrell@sanger.ac.uk and nhl@sanger.ac.uk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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                                                                                                                                                                                                                                                                                                Hall, N., Bowman, S., Lennard, N.J., Doggett, J., Atkin, R., Chillingworth, C., Ormond, D., Harris, B., El-Sayed, N., Hou, L., Melville, S.E., Rajandream, M.A. and Barrell, B.G.
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Trypanosoma brucei
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AL481612.1 GI:11847838
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                                                                                                                                                                                                                                                                                  Direct Submission
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
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/strain="C57BL/6J"
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o. 7.7e+05;
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TITLE
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Best Local :
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13 ACACGTTCTGC 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AA916934 19 bp mRNA linear EST 17-JUN-19 on14a09.sl NCI CGAP Lu5 Homo sapiens cDNA clone IMAGE:1556632 3' similar to SW:BI3_MOUSE P28662 BRAIN PROTEIN I3 ;, mRNA sequence. AA916934
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: nelsayed@tigr.org
Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger Centre are available
at http://www.sanger.ac.uk/Projects/T_brucei/.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seq primer: -40ml3 fwd. ET from Amersham High quality sequence stop: 1.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CDNA Library Preparation: M. Bento Soares, Ph.D. CDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tumor Gene Index
Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy
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                       /lab_host="DH10B"
/clone lib="NCI CGAP Lu5"
/clone lib="NCI CGAP Lu5"
/clone lib="NCI CGAP Lu5"
/note="Organ: lung; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; lst strand cDNA was prepared from neuroendocrine lung carcinoid, and was then primed with a Not I - oligo(dT) primer. Double-stranded CDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library is normalized. Library was
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constructed by Bento Soares and M. Fatima Bonaldo.
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/mol type="genomic DNA"
/strain="TREU927"
/db_xref="taxon:5691"
                                                                                                                                                                                                                                                                                                                   /db_xref="taxon:9606"
/clone="IMAGE:1556632"
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                                                                                                                                                                                                                                                                                        tissue_type="carcinoid"
                                                                                                                                                                                                                                                                                                                                                                              organism="Homo sapiens"
|mol_type="mRNA"
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00.0%; Pred. No.
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AZ820818
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Best Local S
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0093 row: I column: 18
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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2M0093I18F Mouse 10kb plasmid UUGC1M library Mus musculus
clone UUGC2M0093I18 F, genomic survey sequence.
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Location/Qualifiers
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Fax: 801 585 7177
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           /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone_lib="Mouse 10kb plasmid_UUGCIM_library"
/note="Wector: PMP42Nry, Purified_genomic_DNA_from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
(http://www_Jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD2 (gi|473211|gb|AF1290721), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coli XL10-Gold (Stratagene) cells
and selected for ampitillin resistance "
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and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
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100.0%; Pred. No. 2.6e+06;
vative 0; Mismatches 0;
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Contact: Anderson SI
Genomics and Bioinformatics
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             Eukaryota; Metazoa;
Mammalia; Eutheria;
1 (bases 1 to 20)
                                                                                                                                                              AZ780308 20 bp DNA linear GSS 16 2M0017J05R Mouse 10kb plasmid UUGC1M library Mus musculus clone UUGC2M0017J05 R, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Roslin, Midlothian, EH25 9PS, UNITED KINGDOM
Single pass sequencing. Bases called and trimmed with phred
v0.020425.c. Vector identified by cross_match with the -minscore 20
and -minmatch 12 options. Vector:pBlueScriptII(KS+) R. Site 1:
ECORI R. Site 2: NotI Description: Normalised library constructed
from pooled tissue from day 30 placentas. Clones available from UK
Centre for Functional Genomics in Farm Animals, Roslin Institite,
Roslin, Midlothian, UK, EH25 9PS, www.arkgenomics.org.
Location/Qualifiers
                                                                                                                               AZ780308
AZ780308.1 GI:12911838
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EST.
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AJ666323 CSEQRAN09 Sus scrofa cDNA clone C0000033_J06, mRNA
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                                                                       Mus musculus
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                                                                                        Mus musculus (house mouse)
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Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae; Sus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone="C0000033_J06"
/tissue_type="placenta"
/tissue_type="placenta"
/clone_Tib="CSEQRAN09"
/note="Vector: pBlueScriptII(KS+); Site_1: EcoRI; Site_2:
/notI; Single pass sequencing. Normalised library
constructed from pooled tissue from day 30 placentas."
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/db_xref="taxon:9823"
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100.0%; Pred. No. 2.0
1ve 0; Mismatches
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                                  Chordata;
Rodentia;
 Barber, M.,
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                                Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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3. 2.6e+06;
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Beacorn, T.,
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hes 0;
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Duval, B.,
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Mus musculus Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Euteheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 21) [hamil, C., Duval, B., Hamil, C., Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: ddunn@genetics.utah.edu
Insert Length: 1000 Std Error: 0.00
Plate: 0017 row: J column: 05
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
                                                                                                                                                                                      AZ333915
AZ333915.1 GI:10400726
                                                                                                                                                                                                                                   1M0063002F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0063002 F, genomic survey sequence.
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Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   plasmid inserts
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                                                                                                                                                                                                                                                                                            AZ333915
                                                                                                                                       Mus musculus (house mouse)
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801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi |4732114 |gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent B. coli XI10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/clome_lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: pWp42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jack
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC2M0017J05"
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100.0%; Pred. No. 2.6e+06;
1ve 0; Mismatches 0;
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Unpublished (2000)
Contact: Robert B.
                                                                                                                                                                                                                      AZ421118
21 bp JNA
1M0199D16F Mouse 10kb plasmid UUGCIM library Mus
clone UUGC1M0199D16 F, genomic survey sequence
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High quality sequence stop:
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Insert Length: 10000 Std Eror:
Plate: 0063 row: O column: 02
Seq primer: CGTTGTANAACGACGGCCAGT
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Fax: 801 585 7177
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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Mouse whole genome scaffolding with paired end reads from 10kb
               Dunn, D.,
             Mammalia; Eutheria; Rodentia; ()
1 (bases 1 to 21)
Dunn,D., Aoyagi,A., Barber,M.,
                                                                                                                Mus musculus
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                                                                                   Eukaryota;
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/strain="C57BL/6J"
/db_xref="taxon:10090"
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                                                            Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Sutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                            GI:10545131
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               Beacorn, T., Duval, B., Hamil, C.,
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Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                    AI035419

22 bp mRNA linear EST ub46605.rl Soares mammary gland NbMMG Mus musculus CDNA IMAGE.1380777 5' similar to TR.016247 Q16247 HISTONE HITRANSCRIPTION FACTOR LARGE SUBUNIT 2A. ;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0199 row: D column: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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Nouse, whole genome scaffolding with paired end reads from 10kb
                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            High quality sequence stop: 21.
Location/Qualifiers
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                                                                       Mus musculus
                                                                                                                                          AI035419.1
                                                                                                                                                                      AI035419
                                                                                           Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
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(bases 1 to 22)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gi | 4732114 | gb | AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 purified. The sheared, adaptored mean adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /lab_host="E. Coli strain XL10-Gold, T1-resistant, P-"/Clone_lib="Mouse 10kb plasmid UUGCIM library"/note="Vector: pW042nv, Purified genomic DNA from M.musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Laboratory Mouse DNA Resource
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
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100.0%; Pred. No. 2.
ive 0; Mismatches
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REFERENCE
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                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 22
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JOURNAL
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                                                                                          Eukaryota; Metazoa; Chordata; C
Mammalia; Eutheria; Rodentia; S
1 (bases 1 to 23)
Dunn, D., Aoyagi, A., Barber, M.,
                                                                                                                                                                                                                                                                                                          AZ810074 23 bp DNA linear GSS 20-FEB-200200074J19F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC2M0074J19 F, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Trace considered overall poor quality Possible reversed clone: similarity on wrong strand Seq primer: -28ml3 rev2 ET from Amersham High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     rax: 314 200 2007
Email: mouseesfewatson.wustl.edu
This clone is available royalty-free through LLNL; contact the
This clone is available royalty-free through LLNL; contact the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WashU-HHMI Mouse EST Project
Washington University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Marra,M., Hillier,L., Allen,M., Bowles,M., Dietrich,N., Dubu
Geisel,S., Kucaba,T., Lacy,M., Le,M., Martin,J., Morris,M.,
Schellenberg,K., Steptce,M., Tan,F., Underwood,K., Moore,B.,
Theising,B., Wylie,T., Lennon,G., Soares,B., Wilson,R. and
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                           clone UUGC2M0074J19
AZ810074
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (1996)
                                                                                                                                                                                            Mus musculus
                                                                                                                                                                                                                                                                   AZ810074.1 GI:12976974
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IMAGE Consortium (info@image.llnl.gov)
MGI:903245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Marra M/Mouse EST Project
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314 286 1810
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the Not I and Eco RI sites of the modified pT7T3 vector.
RNA provided by Dr. Minoru Ko, Wayne State Univ. Library
constructed and normalized by Bento Soares and M.Fatima
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        T 3'); double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /tissue_type="mammary gland"
//dev_stage="4 weeks"
//lab host="UPH10B"
//clone_lib="Soares_mammary_gland_NDMMG"
/note="Organ: mammary_gland; Vector: pT7T3D-Pac
(Pharmacia) with a modified polylinker; Site 1:
Site_2: Eco RI; 1st_strand_cDNA was primed_with
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100.0%; Pred. No.
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                                                                                                                                             Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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                                                                                               Beacorn, T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 22;
                                                                                                    Duval, B., Hamil, C.,
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                                                                                                                      Mus musculus

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus

1 (bases 1 to 24)
     Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                AZ586137. Q1 bp DNA line 1M0391M17R Mouse 10kb plasmid UUGCIM library Mu clone UUGCIM0391M17 R, genomic survey sequence. AZ586137.1 GI:11708315
                                                                                                                                                                                                                                                                                SSD
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Insert Length: 10000 Std Error: 0.00
Plate: 0074 row: J column: 19
Seq primer: CGTTGTAAAACGACGGCCAGT
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Unpublished (2000)
Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
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801 585 7177
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/strain="C57BL/6J"
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Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome C
University of Utah
Rm. 308, Biomedical Polymer
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Welss,R. Mouse whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                                                                                           AZ829971 24 bp DNA linear GSS 20-FEB-200
2M0107K20R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC2M0107K20 R, genomic survey sequence.
                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 24)
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Plate: 0391 row: M column: 17
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Fax: 801 585 7177
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/strain="C57BL/6J"
/db_xref="taxon:1000"
/clone="UUGC1M0391M17"
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100.0%; Pred. No. 2.6e+06;
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                                          Hall, N., Bowman, S., Lennard, N.J., Doggett, J., Atkin, R. Chillingworth, C., Ormond, D., Harris, B., El-Sayed, N., Helville, S.E., Rajandream, M.A. and Barrell, B.G.
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T. brucei sheared genomic DNA
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Seq primer: CACACAGGAAACAGCTATGACC
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
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Unpublished (2000)
Contact: Robert B.
Direct Submission
Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing
                                                                                                                                                                    Trypanosoma
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Fax: 801 585 7177
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/mol type="genomic DNA"
/strain="C57BL/6J"
/db xref="taxon:10090"
/clone="UUGC2M0107K20"
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                                                                                                                                                                                                                                                                                                                                                                               Tel: 650 723 2227
Fax: 650 725 8221
Email: walbot@stenford.edu
Very probable ligation site of ends cut by single endonuclease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Maize genomic sequences found using engineered 
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Constructed at the Institute for Genomic Research (TIGR),
Rockville, MD. Genomic DNA isolated from a cloned population of
Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically sheared
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                                                                                                                                                                                                                                                                                                            Class: transposon-tagged
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 855 California Ave,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Stanford University
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/clone lib="1119 - RescueMu Grid AA"
/note="Organ: leaf; Vector: RescueMu (engineered from
pBlueScript backbone); Site_1: BamHI; Site_2: BglII;
                                                                             /tissue_type="leaf"
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/lab_host="DH10B"
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/cultivar="mixed background W23/A188/B73/K55"
/db_xref="taxon:4577"
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/mol_type="genomic DNA"
/strain="TREU927"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                   organism="Zea mays"
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/clone="201f10"
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Pred. No.
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RescueMu Grid
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                                                                                                                                                                                                                                                                                                                                                        sequence from source
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                                                                                                                                                                                                                                                                                                                                                        sequence.
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AZ623015/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 ACATGCATCA 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0460 row: F column: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   plasmid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus
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GSS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1M0460F14F Mouse 10kb plasmid UUGC1M library Mu clone UUGC1M0460F14 F, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mus musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            quality sequence stop: 26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 801 585 5606
801 585 7177
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                       /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/clone lib="Mouse 10kb plasmid UUGC1M library"/note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackso. Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                              /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                  /db_xref="taxon:10090"
/clone="UUGC1M0460F14"
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            bp DNA linear GSS 13-DEC-200
UUGC1M library Mus musculus genomic
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o. 2.6e+06;
o;
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                                                                                                                                                                                    Jackson
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SOURCE
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VERSION
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AZ825865/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Class: plasmid ends
High quality sequence stop: |
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0101 row: J column: 14
Seq primer: CGTTGTAAAACGACGGCCAGT
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Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2M0101J14F Mouse 10kb plasmid \bar{t} clone UUGC2M0101J14 F, genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               University of Utah Genome
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus (house mouse)
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                           /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PWP42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackso
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Biomedical Polymers Research Bldg.,
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                                                                                                                                                    aboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                 'sex="Male"
                                                                                                                                                                                                                                                                                                     /db_xref="taxon:10090"
/clone="UUGC2M0101J14"
                                                                                                                                                                                                                                                                                                                                                    /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                    organism="Mus musculus"
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Pred. No.
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smid UUGC1M library Mus musculus genomic
nomic survey sequence.
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2.6e+06;
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                                                                                                                                                                             ic DNA from M. from the Jackson
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adaptored DNA was purified and size-selected 10.5 kb range using preparative agarose gel

for a 9.5 to

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CG723368/c
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ORGANISM
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ORIGIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      88 TGGATGGAGC 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Zea mays
Zea mays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WES California Ave, Palo Alto, CA 94304, USA Tel: 650 723 2227 Fax: 650 723 2227 Email: walbot@stanford.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD clade; Panicoideae; Andropogoneae; Zea.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              survey sequence.
CG723368
CG723368.1 GI:37759145
GSS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Reverse complemented post-ligation sequence from source sequence. Plate: 1119076 row: A column: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Walbot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Maize genomic sequences found using engineered Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Walbot, V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CG723368
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Very probable ligation site of ends cut by single endonuclease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Department of Biological Sciences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TGGATGGAGC 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cord University
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                                /clone_lib="119 - RescueMu Grid AA"
/clone="Organ: leaf; Vector: RescueMu (engineered from /note="Organ: leaf; Vector: RescueMu (engineered from pBlueScript backbone); Site_1: BamHI; Site_2: BgIII;
RescueMu is a 4.9 kb, modified maize Mu transposon designed to allow plasmid rescue from total genomic DNA.
Mu elements insert preferentially into transcription units. For more information on RescueMu, go to the web site 'www.zmdb.iastate.edu' and follow the links for 'RescueMu.' Grid AA was grown at UC San Diego in 2002. DNA was extracted from leaf strips, double digested using BamHI and BgIII, and ligated to form circular plasmids. DH10B cells were transformed and then screened on LB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      transposon-tagged.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10.5 kb range using preparative agarose gel electrophoreeis. Vector DNA was prepared from a derivative of pwD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                   plates with ampicillin."
                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Zea mays"
/mol_type="genomic D
                                                                                                                                                                                                                                                                                                                                 /lab
                                                                                                                                                                                                                                                                                                                                                                           /db_xref="taxon:4577"
/tissue_type="leaf"
                                                                                                                                                                                                                                                                                                                                                                                                                          cultivar="mixed background W23/A188/B73/K55"
                                                                                                                                                                                                                                                                                                                                                   dev_stage="adult"
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                                                                                                                                                                                                                                                                                                                               host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                   _type="genomic DNA"
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Pred. No.
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VERSION
KEYWORDS
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AA636083
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ORGANISM
                                                              DEFINITION
                                                                                 LOCUS
                                                                                                  AI930845/c
                                                                                                                       RESULT 31
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Best Local :
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Best Local Similarity 100.0%;
Matches 10; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              JOURNAL
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                                                                                                                                                                                                                                                                                    Local
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                                                                                                                                                                                                                          39 GAGTTTGCAG 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12 CCCTGCATGC 3
                                                                                                                                                                                      σ
                 AI93008.71 ilnear EST 12 bb mRNA linear EST 12 bb43b01.71 Gm-c1015 Glycine max cDNA clone GENOME SYSTEMS Gm-c1015-2 5' similar to TR:02299 022999 PUTATIVE NAM/NO MERISTEM PROTEIN. ;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AA636083 28 bp mRNA linear EST 31-OC nr15b06.s1 NCI_CGAP_Ew1 Homo sapiens cDNA clone IMAGE:1168019 similar to wP:W02B12.10 CE03770 ;, mRNA sequence.
AI930845
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Trace considered overall poor quality Insert Length: 1170 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tumor Gene Index
Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates; 1 (bases 1 to 28)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Seq primer: -40ml3 fwd. ET from Amersham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tissue Procurement: Lee Helman, M.D., Michael R. Emmert-Buck, M.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AA636083.1 GI:2559022
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          found
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                                                                                                                                                                                                                                                                   Similarity 100.0%; I
10; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cDNA Library Preparation: David B. Krizman, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.B. Consortium/LLNL at:
                                                                                                                                                                                      GAGTTTGCAG 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CCCTGCATGC 58
                                                                                                                                                                                                                                                                                                                                                                                  /clone_lib="NCI_CGAP_Ew1"
/note="Vector: pAWP10; mRNA made from Ewing's sarcoma,
/note="Vector: pAWP10; mRNA made from Ewing's sarcoma,
cDNA made by oligo-dT priming. Non-directionally cloned.
Size-selected on agarose ge1, average insert size 600 bp
Reference: Krizman et al. (1996) Cancer Research
                                                                                                                                                                                                                                                                                                                                                                       56:5380-5383."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone="IMAGE:1168019"
/tissue_type="Ewing's sarcoma"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    organism="Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7.6%; Score 10; .DB 9; Lei
100.0%; Pred. No. 2.6e+06;
                                                                                                                                                                                                                                                                 7.6%; Score 10; DB
100.0%; Pred. No. 2.
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                                                                                                                                                                                                                                                                 2.6e+06;
hes 0;
                                                                                                                                                                                                                                                                                                         DB 1;
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                                        NAM/NO APICAL
                                                           EST 12-JUL-2004
YSTEMS CLONE ID:
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KEYWORDS
                                                      DEFINITION
                                                                                                    AJ807731/c
     ACCESSION
                                                                                                                                  RESULT 32
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Best Local
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                                                                                                                                                                                                              24
AJ807731 Antirrhinum majus whole plant 018 \stackrel{6}{=} 07-j14, mRNA sequence. AJ807731
                                                                                                                                                                                                                                                                  N
                                                                                                                                                                                                                                                                                                                      10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Shoemaker, R., Keim, P., Vodkin, L., Erpelding, J., Coryell, V., Khanna, A., Bolla, B., Marra, M., Hillier, L., Kucaba, T., Martin, J., Beck, C., Wylie, T., Underwood, K., Steptoe, M., Theising, B., Allen, M., Bowers, Y., Person, B., Swaller, T., Gibbons, M., Pape, D., Harvey, N., Schurk, R., Ritter, E., Kohn, S., Shin, T., Jackson, Y., Cardenas, M., McCann, R., Waterston, R. and Wilson, R.
                                                                              AJ807731
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: est@watson.wustl.edu
When it has been determined, an EST from the other end of this
clone is listed in the 'Other ESTs on clone' field. Trace
considered overall poor quality Possible reversed clone: similarity
on wrong strand This clone is available through: Biogenetic
Services, 801 32nd Ave. Brookings, SD 57006 USA (phone: 800 423
4163; email: info@biogeneticservices.com)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Shoemaker R/Public Soybean EST Project Public Soybean EST Project Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. Louis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids I; Fabales; Fabaceae; Papilionoideae; Phaseoleae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AI930845.1 GI:566680
EST.
Glycine max (soybean)
Glycine max
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Fax: 314 286 1810
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                                                                                                                                                                                                                                                                                                                                           Similarity
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                                                                                                                                                                                                              CATGCATCAT 15
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone lib="Gm-cl015"
//clone lib="Gm-cl015"
//clone lib="Gm-cl015"
//clone lib="Gm-cl015"
//clone lib="Gm-cl015"
//clone lib="Gm-cl015"
//clone "Vector: pBluescript II XR; Site 1: EcoRI; Site 2:
XhoI; This cDNA library was constructed from mRNA isolated
from mature flowers of field grown plants. The cDNA
library was prepared using the Stratagene pBluescript II
XR cDNA library construction kit. Complementary DNA was
synthesized from mRNA using a primer consisting of a poly
(dT) sequence with a XhoI restriction site. EcoRI adapters
were ligated to the blunt-ended cDNA fragments followed by
XhoI digestion. The cDNA fragments were directionally
cloned into the EcoRI-XhoI restriction site of the
pBluescript vector. The ligated cDNA fragments were
transformed into XLIO-Gold host cells. This library was
                                                                                                                                                                                                                                                                                                                                                                                                                                             constructed by Dr. Randy Shoemaker and Dr. Erpelding."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /db xref="taxon:3847"
/clone="GENOME SYSTEMS CLONE ID: Gm-c1015-2"
/tissue_type="Mature flowers, field grown plants"
/lab host="XI10-Gold"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Glycine max"
/mol_type="mRNA"
/cultivar="Williams 82"
                                                                                                                                                                                                                                                                                                           100.0%; **
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                                                                                                                                                                                                                                                                                                             Score 10; DB 1; Le
; Pred. No. 2.6e+06;
0; Mismatches 0;
                                                                         28 bp
                                                Antirrhinum
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                                                                         linear
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VERSION
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CF328209
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AUTHORS
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JOURNAL
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1 (bases 1 to 28)

Zachgo, S., Stueber, K., Saedl
Antirrhinum EST collection
Unpublished (2003)
Contact: Schwarz-Sommer Z
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              39 GAGTTTGCAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta;
Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae;
Spermatroideae, Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CF328209 28 bp mRNA linear EST 18-AUG NACL--03-A03.bl Rice callus plasmid cDNA library (NACL) Oryza sativa (japonica cultivar-group) cDNA clone NACL--03-A03, mRNA
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H.,

Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)
                                                                                                                                                                                                                                                                    Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GF328209
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Carl-von-Linne Weg 10, D-50829,
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                           of Bioscience and Bioinformatics,
                                                                                                                                                                                                                                                                                                                                               Genomics and Genetics Institute, GreenGene Biotech Inc.; Division
                                                                                                                                                                                                                                                                                                                                                                      Contact: Nahm B.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CF328209.1
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EST.
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                                                                                                                                                                                                                                                 bhnahm@ggbio.com,
/db xref="Ricaxon:39947"
/db xref="NACL--03-A03"
/tissue_type="callus"
/dev_stage="proliferated callus on 2N6 media for 3
/lab host="E.coli DH10B"
/clone_lib="Rice callus plasmid cDNA library (NACL
/note="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA was
                                                                                                                                                 /mol_type="mRNA"
/cultivar="Nackdong"
                                                                                                                                                                           organism="Oryza sativa (japonica cultivar-group)
/m~l +-ma-"m¤NNN"
                                                                                                                                                                                                                             location/Qualifiers
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/tissue_type="whole plant"
/clone_lib="Antirrhinum ma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mol_type="mRNA"
/db_xref="taxon:4151"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      7.6%;
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Pred. No.
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                                                                                                                                                                                                                                               bhnahm@bio.myongji.ac.kr.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1; L
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                                                                                                                                                                                                                                                                                                                             MyongJi University
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                                                                   for 30 days"
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      capped
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Query Match
Best Local Similarity
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Best Local (
                                                                                                                                                                                                                                                                                                                     misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AL Submitted (23)—CCT-2003) Balzergue S., UMRGV, INRA/CNRS, 2 rue Gaston Cremieux, 91057 Evry cedex, FRANCE PCR was performed on DNA from transformants of Arabidopsis thaliana plants from NRA (versailles). The DNA fragment(s) resulting from the PCR were directly sequenced from the left or the right border to determine the genomic sequence flanking the insertion. T-DNA derived sequences were removed. Information to order the corresponding mutant line and a link to a database providing a graphical display of the insertion site are available at http://dbsgap.versailles.inra.fr/publiclines/. This sequence has been generated in the framework of the French plant genomics program 'Genoplante' (http://www.genoplante.com and http://genoplante-info.infobiogen.fr).

Location/Qualifiers
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                                                                                                    55
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                                                                                                                                                       10;
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Balzergue, S.
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EMBO Rep. 3 (12), 1152-1157 (2002)
22363535
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T-DNA integration into the Arabidopsis genome depends on sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Arabidopsis thaliana (thale cress)
Arabidopsis thaliana
Arabidopsis thaliana
Eukaryota, Viridiplantae, Streptophyta, Embryophyta; Tracheophyta;
Spermatophyta, Magnoliophyta, eudicotyledons, core eudicots;
Spermatophyta, Magnoliophyta, eudicotyledons, core eudicots;
rosids, eurosids II; Brassicales, Brassicaceae, Arabidopsis.
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GSS; left border; T-DNA
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Arabidopsis thaliana T-DNA flanking sequence,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Direct Submission
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                                                       ATGCACACGT 1
                                                                                                  ATGCACACGT 64
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                                                                                                                                                                                                                                                                        /note="T-DNA flanking sequence
left border"
                                                                                                                                                                                                                                                                                                                     /clone_lib="Arabidopsis thaliana T-DNA insertion lines"
1. .29
                                                                                                                                                                                                                                                                                                                                                                                                                organism="Arabidopsis thaliana"
|mol_type="genomic DNA"
|cultivar="Wassillewskija"
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                                                                                                                                                                                                                                                                                                                                                                          clone="356H08"
                                                                                                                                                                           100.0%;
                                                                                                                                                                      7.6%; Score 10;
.00.0%; Pred. No.
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                                                                                                                                       DB 9; Lu., 2.6e+06; 0;
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                                                                                                                                               Gaps
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BE385582/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16 CTACCTGCCG
                                                                                                                                                                                                                                                                                               30 bp DNA linear GSS 04-OCT-201 IM0251A18F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0251A18 F, genomic survey sequence.
AZ451718 AZ451718.1 GI:10607803
GSS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10;
                                          Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rokes, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. Mouse, whole genome scaffolding with paired end reads from 10kb Mouse, whole genome scaffolding with paired end reads.
                                                                                                                                                                 Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
1 (bases 1 to 30)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cDNA Library Preparation: Ling Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.gov Plate: LLCM283 row: e_column: 17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens (human)
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Unpublished (2000)
                                                                                                                                                                                                                                            Mus musculus
                                                                                                                                                                                                                                                                Mus musculus (house mouse)
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National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 30)
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/clone lib="NIH MCC_20"
/note="Organ: skin; Vector: pOTB7; Site 1: XhoI; Site 2: EcoRI; cDNA made by oligo-dT priming. Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies)."
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/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:3616888"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /tissue_type="melanotic melanoma"
/lab_host="DH10B (phage-resistant)"
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00.0%; Pred. No.
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JOURNAL

JOURNAL MEDLINE PUBMED

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REFERENCE AUTHORS

RESULT 34 AJ588718/c

ACCESSION DEFINITION

VERSION **EYWORDS**

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University of Utah Ge
University of Utah
Rm. 308, Biomedical I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19 GACCTGCTGC 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10;
                                                                                                                                                                                                                                                                                                                                                                30 bp DNA linear GSS 16-FEB-20020053K07R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC2M0053K07 R, genomic survey secuence
                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (baees 1 to 30)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Class: plasmid ends
High quality sequence stop: 30.
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Insert Length: 10000 Std Erro
Plate: 0251 row: A column: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seq primer: CGTTGTAAAACGACGGCCAGT
                                               Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                   Mus musculus
                                                                                                                                                                                                                                                                                                                                AZ797441.1 GI:12946521
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                     asmid inserts
                                                                                                                                                                                                                                                                              musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   http://www.jax.org/resources/documents/dnares/). The DNA (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pula? (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid Rl. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/clone_lib="Mouse_10kb_plasmid_UUGC1M_library"
/note="Vector: PW942nv; Purified_genomic_DNA_from_M.
musculus_C57BL/6J_(male)_was_obtained_from_the_Jackso
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mol_type="genomic DNA"
strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 <u>,</u>
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ches 0;
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AZ844017
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Insert Length: 10000 Std Err
Plate: 0053 row: K column: 0
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Fax: 801 585 7177
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University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                  Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 30)
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                       Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
                                                                                                                                                                                                                                                              Mus musculus
                                                                                                                                                                                                                                                                                                                                     AZ844017.1 GI:13013925
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gi|4732114|gb|AFT29072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/clone lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
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Pred. No.
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RESULT 39
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Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19 GACCTGCTGC 28
                                                                                                                                                                                                                                                                                                                                                     CL670968 9 bp DNA linear GSS 09-JUL-200 PRIO163c_B10 - PRIO163c.B21 (9) Mixed stage fosmid library of P. pacificus var. California Pristionchus pacificus genomic, genomic
                                                                                                                                                                                                                                                                                        survey sequence.
CL670968
CL670968.1 GI:50169457
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                                                             Nucleic Acids Res. 32 (1), D421-D422 (2004)
                                                                                Srinivasan,J., Otto,G.W.,
AppaDB: an AcedB database
Pristionchus pacificus
                                                                                                                                                                           Eukaryota; Metazoa; Nematoda; Chromadorea; Diplogasterida; Neodiplogasteridae; Pristionchus.
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Pristionchus pacificus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0143 row: O column: 01
Steq primer: CGTTGTAAAACGACGGCCAGT
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University of Utah Genome Center
University of Utah
                  Evolutionary Biology
                                            Contact: Sommer RJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Class: plasmid ends
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                                                                                                                                                      (bases 1 to 9)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (http://www.jax.org/, resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi 4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone_lib="Mouse 10kb plasmid UUGC1M library" /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
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/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Single pass sequencing. Bases called and trimmed with phred v0.020425.c. Vector identified by cross match with the -minscore 20 v0.020425.c. Vector identified by cross match with the -minscore 20 v0.015. VectoripBlueScriptII(SK+) R. Site1: EccRI R. Site2: NotI 5' Seq Primer M13F Normalised library constructed from pooled early embryos, from 8- cell stage to blastcoysts. Clones available from UK Centre for Functional Genomics in Farm Animals, Roslin Institite, Roslin, Midlothian, UK, EH25 9PS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Anderson, S.I., Finlayson, H.A. and Archibald, A.L. 
Development of CDNA and EST resources for studying reproduction and 
embryo development in pigs and cattle 
Unpublished (2004)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae; Sus.
1 (bases 1 to 11)
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This library was generated at Caltech, Pasadena, USA and end
sequenced at Vancouver, Canada.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Max Planck-Institute for Developmental Biology
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Tel: 00497071601371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 www.arkgenomics.org.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Roslin Institute
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                        /organism="GNUS scrofa"

/mol_type="mRNA"

/db xref="taxon:9823"

/clone="C0005190 F15"

/tissue_type="embryo"

/clone_lib="KN277"

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/strain="California"
/db_xref="taxon:54126"
/clone_lib="Mixed_stage_fosmid_library_of_P.
var._California"
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100.0%;

Score 9; Pred. No.

DB 1; Length 11; . 8.9e+06;

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RESULT 42
CF323664
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AUTHORS
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SOURCE
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BH213431
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Best Local
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Arabidopsis thaliana
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnollophyta; eudicotyledons; core eudicots;
rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                             CF323664 16 bp mRNA linear EST 18-AUG-200:
HDN--04-H04.gl OsHDACl-overexpressing transgenic rice lambda phage
cDNA library II (HDN) Oryza sativa (japonica cultivar-group) cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BH213431 II bp DNA linear GSS 24-OCT-200
SALK 009214 Arabidopsis thaliana TDNA insertion lines Arabidopsis
thaliana genomic clone SALK_009214, genomic survey sequence.
  Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
                                                                                        clone HDN--04-H04,
CF323664
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tel: 858 453 4100 x1752 Fax: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Joseph R. Ecker
Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R., Gadrinab,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L., Shinn,P., Zimmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in the
                                                                    CF323664.1 GI:33795589
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Class: TDNA tagged
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Arabidopsis Genome 
Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Arabidopsis thaliana (thale cress)
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                                                                                                                                                                                                                                                                                                                          TTGCAGCCC 51
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                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              /notes TPCR was performed on Arabidopsis thaliana lines each of which contains one or more TDNA insertion elements. The resultant fragment for each line was directly sequenced to determine the genomic sequence at the site of insertion. Details of the protocols used can be found at http://signal.saik.edu/tdna_protocols.html"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /clone="SALK_009214"
/clone_lib="Arabidopsis_thaliana_TDNA_insertion_lines"
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                                                                                                                                                                                                                                                                                                                                                                6.8%; Score 9; DB 8; Ler 100.0%; Pred. No. 8.9e+06; tive 0; Mismatches 0;
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AUTHORS
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AZ458806
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Best Local :
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Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erri
Plate: 0263 row: O column: 1:
                                                                                                                                                                                                                                                        Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weise, R.

Mouse whole genome scaffolding with paired end reads from 10kb
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Le

Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)
                                                                                                                             University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Res
                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AZ458806.1
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
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Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
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                                                                                                           84112,
                                                                                                                                                                                           Contact: Robert B.
                                                                                                                                                                                                                 Unpublished (2000)
                                                                                                                                                                                                                                       plasmid inserts
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                                                                                                           USA
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/mol type="mRNA"
/cultivar="Nackdong"
/db xref="taxon:39947"
/clone="HDN--04-H04"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone lib="OsHDAC1-overexpressing transgenic rice lambda phage cDNA library II (HDN)"
/note="Vector: pBluescript S/", Site 1: EcoRI; Site 2: XhOI; cDNA was inserted into lambda Uni ZAP XR vector at 5' end with EcoRI and 3' end with XhOI site. mENA was derived from rice Histone Deacetylase overexpression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /tissue_type="callus"
/dev_stage="prolifeerated callus on 2N6 media for 2 weeks"
/lab_host="5.coli SOLR"
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100.0%; Pred. No. 8.9e+06;
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                                                                                                                             Polymers Research
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                        0.00
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VERSION
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AZ651177/c
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1M0521C15R Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGCIM0521C15 R, genomic survey sequence.
AZ651177
AZ651177.1 GI:11786406
GSS.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Bukaryota; Metazoa; Chordata; Sciurognathi; Muridae; Murinae; Mus
Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1 (bases 1 to 19)

Nachar M Reacorn.T., Duval,B., Hamil,C.,
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0521 row: C column: 15
                                                                                                                                                                                                                                                                                                                                                                                                                             plasmid inserts
Unpublished (2000)
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                                                                                                                               Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                  University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert B. Weiss
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Laboratory Mouse DNA Resource
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/strain="C57BL/6J"
/db_xref="taxon:10090"
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Best Local
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                                                                                                                                         Contact: Robert B. We
University of Utah Ge
University of Utah
Rm. 308, Biomedical 1
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.

Mouse whole genome scaffolding with paired end reads from 10kb Mouse whole genome scaffolding with paired end reads from 10kb
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Err
Plate: 0048 row: E column: 0
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                                                                     Tel: 801 585 5606
Fax: 801 585 7177
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/strain="C57BL/6J"
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Mouse whole genome scaffolding with paired end reads from 10kb plasmid insexts
                                                                                                                              University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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Email: ddunn@genetics.utah.edu
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Plate: 0280 row: D column: 22
                                                                 Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                         Contact: Robert B.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1 (bases 1 to 19)
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clone UUGC2M0280D22 R, genomic survey sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Laboratory Mouse DNA Resource
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 clone="UUGC2M0048E05"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /db_xref="taxon:10090"
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/strain="C57BL/6J"
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100.0%; Pred. No.
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d UUGC2M library Mus musculus
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AJ589265
                                                               COMMENT
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Direct Submission
Submitted (23-OCT-2003) Balzergue S., UMRGV, INRA/CNRS, 2 rue
Gaston Cremieux, 91057 Evry cedex, FRANCE
PCR was performed on DNA from transformants of Arabidopsis thaliana
plants from INRA (Versailles). The DNA fragment(s) resulting from
the PCR were directly sequenced from the left or the right border
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AJ589265.1 GI:379388887
AJ589265.1 gI:379388887
GSS; right border; T-DNA flanking sequence
                                                                                                                                                                                                                                                                                                                                                   Brunaud, V., Balzergue, S., Dubreucq, B., Aubourg, S., Samson, F., Chauvin, S., Bechtold, N., Cruaud, C., DeRose, R., Pelletier, G., Lepiniec, L., Caboche, M. and Lecharny, A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AJ589265 19 bp DNA linear Arabidopsis thaliana T-DNA flanking sequence, right 547506, genomic survey sequence.
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                                                                                                                                                                    2 (bases 1 to 19)
Balzergue, S.
                                                                                                                                                                                                                                                    of pre-insertion sites
EMBO Rep. 3 (12), 1152-1157 (2002)
22363535
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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Arabidopsis thaliana
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Location/Qualifiers
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/clone lib="Mouse 10kb plasmid UUGCDM library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6U (female) was obtained from the Jackson
Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 and selected for ampicillin resistance."
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/strain="C57BL/6J"
/db_xref="taxon:10090"
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/mol_type="qenomic DNA"
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| 100.0%; Pred. No. 8.
| 100.0%; Mismatches
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b. 8.9e+06;
ches 0; Indels
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                                                                                                                                                                                                                                      Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0327 row: N column: 06
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ende
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 20)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20 bp DNA linear GSS 05-OCT-20
1M0327N06R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0327N06 R, genomic survey sequence
AZ493004
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Incation/Qualifiers
                                                                                                                                                                                                                                                                                                                                      Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. Mouse whole genome scaffolding with paired end reads from 10kb Mouse whole genome scaffolding with paired end reads
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igh quality sequence stop: 20.
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/cultivar="Wassillewskija"
/db_xref="taxon:3702"
/clone="547E06"
Tab_host="E. Coli strain XL10-Gold, T1-resistant, clone_lib="Mouse 10kb plasmid UUGCIM library"
                                                                                                            mol_type="genomic DNA"
strain="C57BL/6J"
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                                                    sex="Male
                                                                         clone="UUGC1M0327N06"
                                                                                            db_xref="taxon:10090"
                                                                                                                                                      organism="Mus musculus"
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100.0%; Pred. No.
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5. 8.9e+06;
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/* /note="Vector: pWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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Contact: Robert B. University of Utah University of Utah Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.

Nouse whole genome scaffolding with paired end reads from 10kb Mouse whole genome scaffolding with paired end reads from 10kb Email: ddunn@genetics.utah.edu Insert Length: 10000 Std Error: Plate: 0017 row: J column: 05 AZ780308 20 bp DNA linear GSS 16-FEB-20 2M0017J05R Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGC2M0017J05 R, genomic survey sequence. Plate: 0017 row: J column: 05 Seq primer: CACACAGGAAACAGCTATGACC Unpublished (2000) Contact: Robert B. Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Butheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1. (bases 1 to 20) Mus musculus (house mouse) Class: plasmid ends plasmid inserts AZ780308.1 GI:12911838 308, 801 585 5606 801 585 7177 USA Biomedical Genome Weiss Polymers Research Bldg., Center 0.00 20 S. 2030 E., SLC, S

quality sequence stop: 20. Location/Qualifiers

/organism="Mus musculus"
/mol_type="genomic DNA"
/strain="C57BL/6J"

/db_xref="taxon:10090" /clone="UUGC2M0017J05"

/sex="Male" /lab_host="E. Coli strain XL10-Gold, T1-resistant, /clone_lib="Mouse 10kb_plasmid_UUGCIM_library"

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Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0087 row: D column: 09
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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Nouse whole genome scaffolding with paired end reads from 10kb
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Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (2000)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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AZ817897.1 GI:12987805
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                                                                                                                                                                                                                                                          quality sequence stop: 20.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Laboratory Mouse DNA Resource
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/lab_host="E. Coli strain XL10-Gold, T1-resistant,
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                                                       sex="Male"
                                                                                     clone="UUGC2M0087D09"
                                                                                                               db_xref="taxon:10090"
                                                                                                                                                                                                  organism="Mus musculus"
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UUGC1M library Mus musculus
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RESULT 50 AZ817897/c

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Matches Query Match Best Local (

REFERENCE AUTHORS

SOURCE ORGANISM

ACCESSION VERSION

KEYWORDS

JOURNAL COMMENT

COMMENT

cell_type="lymphocytes"

TITLE

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                                                                                                                                                                                                                                                                                                                                                                          Submitted (07-JAN-2002) Hong-Seog Park, Korea Research Institute of Bioscience and Biotechnology (KRIBB), Genome Research Center (GRC); 52, Oun-dong, Yusong-gu, Daejeon 305-333, Korea (B-mail:redstone@mail.kribb.re.kr, URL:http://phs.grc.kribb.re.kr/, Tel:82-42-866-7181, Fax:82-42-860-4409)
Clones are derived from the chimpanzee BAC library RP-43 This BAC
                                                                                                                                                                                                                                                                                                                   end was generated during the R&D process and may have higher chance of clone tracking errors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Park,H., Kim,Y., Kim,S., Han,Y., Woo,T., Park,K., Eun,C.J., Hoon,S.T., Chu,M., Kim,H., Joo,S., Kim,C., Song,W. and Yoo,H. Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Park,H., Kim,Y., Kim,S., Han,Y., Woo,T., Park,K., Eun,C.J., Hoon,S.T., Chu,M., Kim,H., Joo,S., Kim,C., Song,W. and Yoo,H. BAC end sequences of Library RP-43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Pan.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AG189452 20 bp DNA linear GSS 06 Pan troglodytes DNA, clone: RP43-063P18.TJ, genomic survey
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/clone="RP43-063P18.TJ"
/sex="male"
                                                                        /organism="Pan troglodytes"
/mol_type="genomic DNA"
                                                                                                                               1. .20
                                                    /mol_type="genomic DN
/db_xref="taxon:9598"
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Roslin, Midlothian, EH25 9PS, UNITED KINGDOM

Single pass sequencing. Bases called and trimmed with phred

Single pass sequencing. Bases called and trimmed with phred

v0.020425.c. Vector identified by cross match with the -minscore 20

and -minmatch 12 options. Vector:pBlueScriptII(KS+) R. Site 1:

BCORI R. Site 2: NoII Description: Normalised library constructed

from pooled tissue from day 30 placentas. Clones available from UK

Centre for Functional Genomics in Farm Animals, Roslin Institite,

Roslin, Midlothian, UK, EH25 9PS, www.arkgenomics.org.

Roslin, Midlothian, UK, EH25 9PS, www.arkgenomics.org.
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                                                                                                       AZ323807 21 bp DNA linear GSS 29-SEP-200100045N21F Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGCIM0045N21 F, genomic survey sequence.
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Development of cDNA and BST resources for studying reproduction and 
embryo development in pigs and cattle 
Unpublished (2004)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae;
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Sus scrofa
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                     Mus musculus
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                    Mus musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Sus scrofa"
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/mol type="mRNA"
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/clone="C0000044_J13"
/tissue_type="placenta"
/clone_Tib="CSEQRANO9"
/note="Vector: pBlueScriptII(KS+); Site_1: EcoRI; Site_2:
NotI; Single pass sequencing. Normalised library
constructed from pooled tissue from day 30 placentas."
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100.0%; Pred. No. 8.9e+06;
tive 0; Mismatches 0;
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Sus.
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                                                                                                                              DEFINITION
                                                                                                                                                                                      RESULT 54
                                                                                              ACCESSION
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Fax: 801 585 7177
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Unpublished (2000)
                     Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert B.
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                           AZ598709.1
                                                                                            AZ598709
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                                                                                                                                                                                                                                                                                                                                     Similarity
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B.; Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R. Mouse whole genome scaffolding with paired end reads from 10kb Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                    1M0413J08R Mouse 10kb plasmid clone UUGC1M0413J08 R, genomic
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Location/Qualifiers
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Insert Length: 10000 Std Error:
Plate: 0045 row: N column: 21
Seq primer: CGTTGTAAAACGACGGCCAGT
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was hunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt end-sified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /lab host="E. Coli strain XL10-Gold, T1-resistant, F-" /clone lib="Mouse 10kb plasmid UUGC1M library" /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mol_type="genomic DNA"
/strain="C57BL/6J"
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                                                           GI:11720899
                                                                                                                                                                                                                                                                                                                                                                                                                                                     6.8%; Score 9; DB 8; Ler 100.0%; Pred. No. 8.9e+06; tive 0; Mismatches 0;
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ACCESSION
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AUTHORS
                                                                 SOURCE
                                                                                                                                                                               DEFINITION
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Best Local Similarity
                                           ORGANISM
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                                                                                                                                         AZ794048 21 bp DNA linear GSS 16-FEB-20
2M0047H07R Mouse 10kb plasmid UUGCIM library Mus musculus genomic
clone UUGC2M0047H07 R, genomic survey sequence.
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                                 Mus musculus
       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                         Mus musculus (house mouse)
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Insert Length: 10000 Std Error: 0.00
Plate: 0413 row: J column: 08
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                      GGGCTGGAT 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD42 (gi|4732114|gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
chemically-competent E. coli XII0-Gold (Stratagene) cells
and selected for ampicillin resistance."
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/strain="C57BL/6J"
/db_xref="taxon:10090"
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/clone_lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
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                                                                                                       GI:12939619
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100.0%; Pred. No. 8.9e+06;
ive 0; Mismatches 0; Indels
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                            Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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Query Match
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GSS.
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Mus musculus (house mouse)
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Dunn, D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T.,
Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von
Niederhausern,A. and Wright,D., Weiss,R.
Mouse whole genome scaffolding with paired end reads from 10kb
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Location/Qualifiers
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Seq primer: CACACAGGAAACAGCTATGACC
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Insert Length: 10000 Std Error:
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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Fax: 801 585 7177
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/strain="C57BL/6J"
/db_xref="taxon:10090"
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2M0105C05R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC2M0105C05 R, genomic survey sequence.
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Insert Length: 10000 Std Error: 0
Plate: 0051 row: A column: 10
Seq primer: CACACAGGAAACAGCTATGACC
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Fax: 801 585 7177
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                    Mus musculus
                                                       Mus musculus (house mouse)
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
/note="Vector: PW042nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackso
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/strain="C57BL/6J"
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Fax: 801 585 7177
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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Plate: 0105 row: C column: 05
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/strain="C57BL/6J"
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(house mouse)
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                                                                                                                                                                                                                                                                                                                                             16 TGCGTGAGT
                                                                                            sequence.
AG189469
AG189469.1 (
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                           Pan troglodytes (chimpanzee)
Pan troglodytes
                                                                                                                                                                                        Pan troglodytes DNA, clone: RP43-064A05.TJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: ddunn@genetics.utah.edu
Insert Length: 10000 Sdd Error:
Plate: 0116 row: K column: 23
Seq primer: CGTTGTAAAACGACGGCCAGT
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University of Utah Genome Center
University of Utah
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity 100.0%;
9; Conservative
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Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD42 (gi|4732114|gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coll XL10-Gold (Stratagene) cells
and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      musculus C57BL/6J (male) was obtained
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mol_type="genomic DNA"
/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22 bp mRNA linear BST 21-DEC-wi39b04.x1 NCI_CGAP_CO16 Homo sapiens cDNA clone IMAGE:2392591 similar to TR:Q02393 HUMAN PAPILLOMAVIRUS 18 E5 CENTRAL SEQUENCE MOTIF PROTEIN 1 ;, mRNA sequence.
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PRIMERS
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NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute, Cancer Genome Anatomy Pro
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Clones are derived from the chimpanzee BAC library RP-43 This BAC Clones are derived from the chimpanzee BAC library RP-43 This BAC
                                 CDNA Library Preparation: M. Bento Soares, Ph.D. CDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                        Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue_Procurement: Ilan Kirsch, M.D., Michael R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens (human)
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Park,H., Kim,Y., Kim,S., Han,Y., Woo,T., Park,K., Eun,C.J., Hoon,S.T., Chu,M., Kim,H., Joo,S., Kim,C., Song,W. and Yoo,H. Direct Submission
                                                                                                                                                                                                                                                                                     Tumor Gene Index Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AI738818.1 GI:5100799
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Unpublished
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/mol_type="genomic DNA"
/db xref="taxon:9598"
/clone="RP43-064A05.TJ"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /cell_type="1ymphocytes"
/clone_lib="RP-43 Chimpanzee Male BAC Library"
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o. 8.9e+06;
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Sequencing Center information can be

Project (CGAP),

EST 21-DEC-1999

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MPI fuer Zuechtungsforschung
Carl-von-Linne Weg 10, D-50829,
Location/Qualifiers
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Zachgo, S., Stueber, K., Saedler, H.,
Antirrhinum EST collection
Unpublished (2003)
Contact: Schwarz-Sommer Z
                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids; lamiids; Lamiales; Plantaginaceae; Antirrhineae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antirrhinum majus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AJ795973 Antirrhinum majus whole 018 3 09 J22, mRNA sequence. AJ795973 GI:51111301
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Location/Qualifiers
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                                                                                                                                         /clone="018_3_09_j22"
/tissue_type="whole plant"
/clone_lib="Antirrhinum majus whole plant"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Subtraction by Bento Soares and M. Fatima Bonaldo.
                                                                                                                                                                                            /mol_type="mRNA"
/db_xref="taxon:4151"
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/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:2392591"
                                                                                                                                                                                                                          organism="Antirrhinum majus"
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100.0%; Pred. No. 8:
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No. 8.9e+06;
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VERSION
KEYWORDS
SOURCE
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AZ307716
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                            S
                                                                                                                              ORIGIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURES
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                                                               Matches
                                                                             Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE
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12
                               50
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University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AZ307716.1 GI:10346985
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus (house mouse)
                                                                             Similarity
CCTGCATGC 20
                              CCTGCATGC 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ass: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (bases 1 to 22)
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801 585 7177
                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sex="Male"
                                                                             6.8%;
100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Weiss
                                                              0
                                                                           Score 9;
Pred. No.
                                                               Mismatches
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22 bp DNA linear GSS 29-SEP-20(
1M0009N02R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0009N02 R, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0009 row: N column: 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hami
Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T.
Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von
Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    High quality sequence stop: 22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mouse whole genome scaffolding with paired end reads
                                                                                                                                                       10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through & 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /lab host="E. Coli strain XI.10-Gold, T1-resistant, F-"
/clone_lib="Mouse 10kb plasmid UUGCIM library"
/notee="Vettor: PWP02nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Biomedical Polymers Research Bldg.,
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/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
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Rodentia;
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DB 8; JG. 8.9e+06; 0;
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AZ416988
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VERSION
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T.,
Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von
Niederhausern, A. and Wright, D., Weiss, R.
Nouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          High quality sequence stop: 22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: ddurn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0192 row: P column: 23
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (2000)
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GSS.
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1M0192P23F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0192P23 F, genomic survey sequence.
                                                                                                                                                              9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus
                                                                                 CAGGACCTG 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ass: plasmid ends
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801 585 7177
                                                                                                                                                                                                                                                                                                                                     Laboratory Mouse DNA Resources (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNMD42 (gil 4732114 | gb| AFT29072 1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/clone_lib="Mouse_10kb_plasmid_UUGCIM_library"/note="Vector: PWD42nv; Purified_genomic_DNA_from_M.musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_f
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/strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 clone="UUGC1M0192P23"
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        22
                                                                                                                                                                                                   6.8%;
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                                                                                                                                                                                                   Score 9; ; Pred. No.
                                                                                                                                                              0;
                                                                                                                                                              Mismatches
                                                                                                                                          ; DB 8; L. No. 8.9e+06; 0;
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                                                                                                                                                                                                                                Length 22;
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High quality sequence stop: 22.
Location/Qualifiers
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Seq primer: CGTTGTAAAACGACGGCCAGT
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Insert Length: 10000 Std Error:
Plate: 0435 row: J column: 11
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Fax: 801 585 7177
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Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pNP42 (gi|4732114|gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coli XI10-Gold (Stratagene) cells
and selected for ampicillin resistance."
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/mol_type="genomic DNA"
/strain="C57BL/6J"
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/clone="UUGC1M0435J11"
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Insert Length: 10000 Std Error:
Plate: 0007 row: D column: 04
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Fax: 801 585 7177
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20
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                                                                                                                                                                                                                           (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity: The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent B. coli XI10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                         Conservative
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/strain="C57BL/6J"
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/clone="UUGC2M0007D04"
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100.0%; Pred. No. 8.
tive 0; Mismatches
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Insert Length: 10000 Std Error:
Plate: 0036 row: 0 column: 22
Seq primer: CGTTGTAAAACGACGGCCAGT
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                                                                                                    adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWP42 (gi|4732114 (gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                   (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /Clone lib="Mouse 10kb plasmid UUGC1M library" /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Biomedical Polymers Research Bldg., 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone="UUGC2M0036022"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    'sex="Male'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               db_xref="taxon:10090"
6.8%; 5c.
100.0%; Pr
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Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Center
Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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10 ATCTGCCAG 18 8 ATCTGCCAG

16

9

Conservative

Similarity

Score 9; Pred. No. Mismatches

Length 22;

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Gaps

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Query Match
Best Local Similarity
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AZ990555
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                                               50 CCTGCATGC 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Musinae; Mus. 1 (bases 1 to 22)

Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R. Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0274 row: N column: 14
Seg primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    High quality sequence stop: 22.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    University of Utah Genome Center University of Utah
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         plasmid
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GSS.
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22 bp DNA linear GSS 27-APR-200200274N14F Mouse 10kb plasmid UUGC2M library Mus musculus genomic clone UUGC2M0274N14 F, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Class: plasmid ends
CCTGCATGC 16
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801 585 7177
                                                                                                     Conservative
                                                                                                                                                                                                                  (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi 4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent B. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/clone_lib="Mouse 10kb plasmid UUGC2M library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (female) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /clone="UUGC2M0274N14"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sex="Female"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         db_xref="taxon:10090"
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                                                                                                                       Score 9; DB 8; Le ; Pred. No. 8.9e+06;
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RESULT 69
BM396286/c
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AW249584/c
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ACCESSION
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RS NIH-MGC http://mgc.nci.nih.gov/.

AL Unpublished (1999)

Other EST9: 2821490.5prime
Contact: Robert Strausberg, Ph.D.

Email: cgapbs-remail.nih.gov

Tissus Procurement: DCTD/DTP cDNA Library Preparation: Ling
Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E.
Consortium (LLNL) DNA Sequencing by: Berkeley MGC sequencing
project Clone distribution: MGC clone distribution information can
be found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html Base Calling / Quality
Scores: PHRED from University of Washingtion Genome Center
Trimming: cross_match from University of Washingtion Genome Center
PHRAP suite. Poly-T Identification: parkatch, pf from Berkeley
Drosophila Genome Project. University of Washingtion Genome Center:
http://www.genome.washington.edu Low Quality Sequence: contiguous
PHRED high quality bases following vector sequence: 9 contiguous
Quality Sequence: Trace file contained 23 contiguous distinct peaks
following vector sequence. Polyadenylation: Based upon the presence
of a XhoI site followed by a run of 14 or more T residues at the
beginning of the sequence, this cDNA insert was polyadenylated.
Plate: LLCM6 row: P column: 3
High quality sequence stop: 9.
Location/Qualifiers

1 23
                                                                                                                                                                                                                                    23
                     23 bp mRNA linear EST 1
5009-0-2-A10.t.2 Chilcoat/Turkewitz cDNA (large fraction)
Tetrahymena thermophila cDNA, mRNA sequence.
   BM396286
                                                                                  BM396286
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2821490.3prime NIH_MGC_7 Homo sapiens cDNA clone IMAGE:2821490
mRNA sequence.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                    Similarity 9; Conserv
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                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /cell lTne="MGC3"
/lab Nost="DH108 (phage-resistant)"
/lab Nost="DH108 (phage-resistant)"
/clone lib="NH1MGC 7"
/clone lib="NH1MGC 7"
/note="Organ: lung; Vector: pOTB7; Site 1: XhoI; Site_2: EcoRI; cDNA made by oligo-dT priming. Directionally cloned into EcoRI/XhoI sites using the following 5; adaptor: GGCACGAG(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies)."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Homo sapiens"
/mol_type="mRNA"
/db xref="taxon:9606"
/clone="MAGE:2821490"
/tissue_type="small cell carcinoma"
/cell_line="MGC3"
                                                                                                                                                                                                                                                                                                                                                                         100.0%;
                                                                                                                                                                                                                                                                                                                                                 6.8%; Score 9; DB 2; Le
100.0%; Pred. No. 8.9e+06;
ive 0; Mismatches 0;
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                                                                                  EST 17-JAN-2002
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                  Pathogen Sequencing Unit
The Sanger Institute The Wellcome Trust Genome Campus
Hinxton, Cambridge, CB10 1SA, UK
Request for clones, please contact: Mike Lehane
prof. M.J.Lehane
School of Biological Sciences,
University of Wales,
Bangor LL57 2UW
All clones with suffix q1c are reverse primer reads st
                                                                                                                                                                                                                                                                                                                                                                                      Glossina morsitans morsitans
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Hippoboscoidea; Glossinidae; Glossina.
                                                                                                                                                                                              14519198
Contact: Hall N
                                                                                                                                                                                                                                    response genes
Genome Biol. 4 (10),
22881942
                                                                                                                                                                                                                                                                                                         Lehane, M.J., Aksoy, S., Gibson, W., Kerhornou, A., Berriman, M., Hamilton, J., Soares, M.B., Bonaldo, M.F., Lehane, S. and Hall, N. Adult midgut expressed sequence tags from the tsetse fly Glossina
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BX558114 Glossina morsitans morsitans adult infected morsitans morsitans cDNA clone Tse36g02_q1c, mRNA sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tel: 773 702 4374
Fax: 773 702 3172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      University of Chicago
920 E. 58th Street, Chicago,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Turkewitz AP
Molecular Genetics and Cell Biology
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Turkewitz, A.P., Karrer, K.M.,
Frankel, J. and Klobutcher, L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tetrahymena thermophila
Eukaryota; Alveolata; Ciliophora; Oligohymenophorea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BM396286.1
EST.
                                                                                                                                                                                                                                                                                             morsitans morsitans and expression analysis of putative immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Glossina morsitans morsitans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BX558114.1 GI:33429261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EST from Tetrahymena thermophila, strain CU428.1, growing cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hymenostomatida; Tetrahymenina; Tetrahymena.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: apturkew@midway.uchicago.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GCGTGAGTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GCGTGAGTT 43
                                                                                                                                                                                                                                                                                                                                                                        (bases 1 to 23)
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clones with suffix qlc are reverse primer reads starting of the cDNA all plc reads are from \,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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/clone lib="Chilcoat/Turkewitz cDNA (large fraction)"
/note="Vector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (2)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /mol_type="mRNA"
/strain="CU428.1"
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100.0%; Pr/
0;
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                                                                                                                                                                                                                                                        R63 (2003)
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. 8.9e+06;
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ORGANISM
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AQ934937
LOCUS
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VERSION
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  Query Match
Best Local S
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: jiewenl@bcn
Plate: 443 row: Comprimer: T7
Seq primer: T7
Class: BAC ends.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tel: 713 798 6514
Fax: 713 798 8142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Baylor College of Medicine
T926, One Baylor Plaza, Ho
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 (bases 1 to 23)
Bradley, A., Li, J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        survey sequence.
AQ934937
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Li, Jiewen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BAC End Sequences from
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Mammalia; Eutheria; Rodentia;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          jiewenl@bcm.tmc.edu
443 row: J column:
                                                                                                                                       /clone lib="RPCI-23"
/note="Organ: Kidney/Brain; Vector: pBACe3.6; Site 1:
/note="Organ: Kidney/Brain; Vector: pBACe3.6; Site 1:
EcoRI; Site 2: EcoRI; Pemale C57BL/6J mouse kidney and/or brain genomic DNA was isolated and partially digested with a combination of EcoRI and EcoRI Methylase. Size selected DNA was cloned into the pBACe3.6 vector at the EcoRI sites. The ligation products were transformed into the pBACe3.6 vector at the pBACe3.6 vector at the sites. The ligation products were transformed into the pBACe3.6 vector at the pBACe3.6 vector at the sites. The ligation products were transformed into the pBACe3.6 vector at the sites. The ligation products were transformed into the pBACe3.6 vector at the second partial vector at t
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/tissue_type="adult infected gu
/clone_lib="Glossina morsitans
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/sub_species="morsitans"
/db_xref="taxon:37546"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 organism="Mus musculus"
/mol_type="genomic DNA"
/strain="C57BL/6J"
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f.brucei"
                                                                                                                                                                                                                                                                                                                                                           'lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                       sex="Female"
                                                                                                                                                                                                                                                                                                                                                                                                                  'clone="443J16-T7"
                                                                                                                                                                                                                                                                                                                                                                                                                                              db_xref="taxon:10090"
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  6.8%; 5c.
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100.0%;
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                                      Score 9; 1
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Pred. No. 8.9e+06;
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DB 8; L. No. 8.9e+06; 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sciurognathi;
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                                                            Length 23;
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9;

Conservative

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Hou, L.,

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: ddunn@genetics.utah.edu
Insert Length: 1000 Std Error: 0.00
Plate: 0150 row: J column: 02
Seq primer: CACACAGGAAACAGCTATGACC
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                    Similarity 9; Conserv
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801 585 7177
                                                                                                                     (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/mol_type="genomic DNA"
/strain="C57BL/6J"
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Constructed at the Institute for Genomic Research (TIGR),
Constructed at the Institute for Genomic Research (TIGR),
Rockville, MD. Genomic DNA isolated from a cloned population of
Rypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically shear
to give a tight size distribution (
4 kb). The v + i method used for the library construction is
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insert libraries for whole genome shotgun sequencing projects.
Genome Sequencing: A Practical Approach, eds. M. Vaudin and B.
Barrell, Oxford University Press, 1999).
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         Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.E., Rajandream,M.A. and Barrell,B.G.
                                                                                                                                                Trypanosoma brucei
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T. brucei sheared genomic genomic survey sequence.
AL466773
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Location/Qualifiers
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/strain="TREU927"
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Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned population of Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically sheared to give a tight size distribution (

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                                                                                                                                                                                                                      L Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and nh@sanger.ac.uk

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hall, N., Bowman, S., Lennard, N.J., Doggett, J., Atkin, R., Chillingworth, C., Ormond, D., Harris, B., El-Sayed, N., Hou, L., Melville, S.E., Rajandream, M.A. and Barrell, B.G.
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T. brucei sheared genomic DNA clone 269e10,
                                                                                                         Details of T. brucei sequencing at the Sanger Centre are available at http://www.sanger.ac.uk/Projects/T_brucei/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Trypanosoma brucei
Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GSS.
Trypanosoma brucei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             genomic survey sequence.
AL484437
AL484437.1 GI:11851207
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Details of T. brucel sequencing at the Sanger Centre are available
at http://www.sanger.ac.uk/Projects/T_brucel/.
Location/Qualifiers
                                                                                                                                                                                                       Email: nelsayed@tigr.org
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CCCTGCATG 9
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/organism="Trypanosoma brucei"
/mol_type="genomic DNA"
/strain="TREU927"
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/mol type="genomic DNA"
/strain="TREU927"
/db xref="texon:5691"
/clone="143d12"
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/db_xref="taxon:5691"
/clone="269e10"
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ORIGIN

Query Match 6.8%; Score 9; DB 9; Length 23;
Best Local Similarity 100.0%; Pred. No. 8.9e+06;
Matches 9; Conservative 0; Mismatches .0; Indels 0; Gaps
Qy 87 CTGGATGGA 95

0

Db 3 CTGGATGGA 11

Search completed: February Job time: 412.351 secs

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2005, 23:32:05

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Maximum
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11170.029 Million cell updates/sec
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Copyright (c) 1993 - 2005 Compugen Ltd
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                AAF30356
ABK56533
ABK56534
ABK56531
ABK56532
ACC68011
ADC59459
AAV64426
ADF74517
AAS98254
AAQ95923
ABK5651593
AAH27321
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Aaf30356 Human Che
Abk55933 Human CLC
Abk56534 Human CLC
Abk56532 Human CLC
Abk56532 Human CLC
Acc68011 Murine ol
Adc59459 Human epi
Aav64426 Pleckstri
Av64426 Pleckstri
Av64426 Pleckstri
Av64427 Forward p
Aas98254 Human pla
Aag95923 Primer A
Abk55934 Human CLC
Abz65159 Human TSG
Abac6159 Human TSG
Ah27321 Human TSG
Ah27375 PCR prime
Aah27375 PCR prime
Aav08740 PCR prime
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ADM622169
ADM76915
AAA29653
AAA29653
AAA296537
AAA719285
AAA296537
AAA77337
AAG70579
AAM77337
AAG70579
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AAM775938
AAM775938
AAM717327
AAM77337
AAM7430
                                            AAX95827

AAC79573

AAF80036

AAH42633

ABK87895

ABX78173

ADF87753

ADF87753

ADF87292972

ABZ992912

ABZ992912

ABZ92912

ABZ92912

ABZ92912

ABZ92912

ABZ92912

ABZ92913

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AAX95935
                                         Aad30197 Human UGT
Abx97303 Human NOV
Abx97303 Human NOV
Abx97303 Human NOV
Adh13368 Human NOV
Adh13368 Human NOV
Adh62212 Human NOV
Adh62213 Hybrid to
Aaa29634 Hybrid to
Aaa2963 Human Mic
Aax19285 HIV-1 gag
Abk51801 Human GDM
Abn02193 Human GDM
Abn02194 Human GDM
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Abx6535 Faat P2X3
Aad497315 Primer of
Add195356 Rat P2X3
Aad497315 Primer us
Abx66975 HCV DNAzy
Adm11727 Primer us
Abx66975 PCR prime
Aax95827 PCR prime
A
  Aaf96648 Human gen
Aah20433 PCR prime
Aac84823 Human TLR
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ABS76660
AAD28830
ABQ82776
ABG68586
ABG683158
ABT09837
                                                               AAL59954
ABQ77009
ACA94795
ABX08586
ABX08586
AAD49565
ABX99250
ACC48666
AAD54826
                   ABT43639
ADA09686
ACF05362
                                              AAL62812
ACD91458
                                                                                                                                                                           AAD42442
AAD46094
AAD30863
                                                                                                                                                                                                                                                                    AAX76633
ABA03087
AAD20034
AAD13315
AAD13214
ABN87318
AAC83865
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AAQ23795
AAQ39379
AAT29187
AAQ86318
AAV41807
AAV41790
AAZ32542
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AAA15505
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ADG77102
ADN11934
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ADJ95469
                                                                                                                                                                                                                                                                                                                                                                                                         Aba315505 Human G-a
Abk93310 PCR prime
Aad47282 Human RT-
Adg77102 V-gene pr
Adn11934 T cucumer
                                                                                                  Aca94795
Abx08586
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Adj95469 Mouse Ubi
Adl95361 Oligoribo
Adl95359 Oligoribo
Adl95358 Oligoribo
Adl95360 Oligoribo
Adl95360 Oligoribo
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         6 Human ant
2 Human VL
53 Human HGP
                                  PCR prime
Anti-HGPR
Human VL
Human GPC
Human GPC
Human ant
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Human TCAN
Human ant
Human TR4
Human TR4
Human TR4
Human Ijg
PCR prime
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Primer Hu
Kappa-cha
HuVk5a ka
Wilson di
Human pan
Human pan
Human pan
PCR prime
Human ant
Human vI
Human VI
Human VI
Human VI
Human VI
Human Ant
Human TI
Human II
Human MC
Wkappa ba
PCR prime
Novel met
Human imm
Human HDG
K+beta M6
Human HGP
K+beta M6
Human HGP
K+beta M6
Human HGP
HUMA
                RESULT 2
ABK55933/c
ABK55933 standard;
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                                                                                                  Query Match
Best Local S
Matches 17
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ABK55933;
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ALIGNMENTS
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02-JUL-2002

(first

entry)

RNA;

17

17; μ

Conservative

0

Mismatches

Indels

0

Gaps

0

ACATGCATCATCTGCCA 17 ACATGCATCATCTGCCA

9

Similarity

Human

CLCA1

gene

enzymatic nucleic acid #304.

tis; cystic fibr bronchodilator;

chronic

obstructive

obstructive pulmonary disease; COPD; a fibrosis; obstructive bowel syndrome; fibrosis; obstructive bowel syndrome; muc

vaccination; mucokinetic;

asthma;

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RESULT 1
AAF30356/c
ID AAF30356 s
XX
XX
AC AAF30356;
XX
DT 14-MAY-200
XX
DE Human chec
XX
KW Checkpoint
KW mitosis;
  Checkpoint with forkhead associated domain and ring mitosis; cell cycle; tumour; diagnosis; antitumour;
                                  Human checkpoint gene chfr 3' PCR primer.
                                                        14-MAY-2001
                                                                                                  standard;
                                                       (first entry)
                                                                                                DNA;
                                                                                                  28
finger; Chfr; human;
drug screening;
                                                                        _
  Homo
                                                              chronic bronchitis; oxygen therapy; bron acetylcysteine.
  14-FEB-2002.
                                                                                                       Human; chloride channel calcium activated 1; CLCA1; ss; antiasthmatic;
                      WO200211674-A2
                                                                                              antiinflammatory;
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The present sequence is that of a 3' PCR primer, used with the 5' primer CC given in AAF30355, to amplify a cDNA fragment corresponding to mucleotides 352-1055 of the human mitotic checkpoint protein Chfr (see Cfir gene encodes the human mitotic checkpoint protein Chfr (see CC AAB20219), which is required for regulation of the transition of cells from prophase to metaphase during mitosis. Loss of expression of Chfr is associated with a predisposition to tumourigenesis upon exposure to mitotic stress. A set of primers (see AAF30353-76) was used to amplify regions spanning the entire chfr coding region in order to determine whether the chfr gene is mutated in any of the human cancer cell lines SW480, DID1, HT29, HCT116, SAOS2, UZOS, IMR5 and NGP. A mutation leading to a Val-580 to Met amino acid substitution was identified in the chfr gene of UZOS cells. Chfr polypeptides and chfr mucleic acids are used in methods of diagnosing tumourigenic cells and of screening for drugs which can inhibit the activity of Chfr in a cancer cell, rendering it more sensitive to additional antitumour therapies
                                                                                                                                                                                                                                                                                                                                                                                                                            Novel nucleic acid sequence of mitotic checkpoint gene encoding a checkpoint with forkhead-associated domain and ring finger protein, diagnosing tumorigenic cells and in screening for anticancer drugs.
                                                    Sequence 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08-FEB-2001.
                                                                                                                                                                                                                                                                                                                                                                                              Example 3; Page 38; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Halazonetis T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-JUL-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ubiquitin-protein ligase; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (WIST-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-JUN-2000; 2000WO-US016391
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                                                    BP; 10 A; 4 C; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99US-0146194P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Scolnick
12.9%;
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                                                    G; 6
 Score 17;
Pred. No.
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                                                    T; 0
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DB 5; L
                                                    U; 0 Other;
              Length
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RESULT 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to enzymatic nucleic acid molecules that down cregulate expression of chloride channel calcium activated 1 (CLCA1) genes by cleaving RNA derived from the genes. The nucleic acid sequences are useful as pharmaceutical agents for treating conditions such as chronic obstructive pulmonary disease (COPD), chronic bronchitis, asthma, cystic fibrosis, obstructive bowel syndrome and any other diseases or conditions that are related to or will respond to the levels of CLCA1 in a cell or tissue. The sequences are useful for reducing CLCA1 activity in a cell, chance, are useful for treatment of a partient having a condition CC associated with the level of CLCA1, where the invention further comprises the use of one or more therapies under conditions suitable for the treatment, for example, oxygen therapy, bronchodilators, corticosteroids, antibacterials, vaccinations, acetylcysteine and mucokinetic agents. The nucleic acids of the invention are also used as diagnostic tools to carymatic most of CLCA1 and mutations within diseased cells or to detect the presence of CLCA1 and mutations within diseased represents an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
                                                                                                                                                               Human; chloride channel calcium activated 1; CICA1; 88; antiasthmatic; antiinflammatory; chronic obstructive pulmonary disease; COPD; asthma; chronic bronchitis; cystic fibrosis; obstructive bowel syndrome; chronic bronchitis; cystic fibrosis; obstructive bowel syndrome; oxygen therapy; bronchodilator; corticosteroid; vaccination; mucokinet.
                                                                                                                                                                                                                                                                                                                                                 533/c
ABK56533 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Enzymatic polynucleotide that down regulates expression of chloride channel calcium activated gene, useful for treating Chronic obstruction pulmonary disease (COPD), chronic bronchitis and asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-AUG-2000; 2000US-0224383P
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                                                                                                                                                                                                                                                   Human CLCA1
                                                                                                                                                                                                                                                                                                                      ABK56533;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 4; Page 58; 152pp; English.
               09-AUG-2001; 2001WO-US024970
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             enzymatic
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                                                                                WO200211674-A2
                                                                                                                Homo sapiens
                                                                                                                                                 acetylcysteine.
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SYNTEX USA LLC.
THOMPSON J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nucleic acid molecule of the invention
                                                                                                                                                                                                                                                  gene enzymatic nucleic acid #904
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                                                                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11.4%; 500
100.0%; Pr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            G; 0 T; 3 U; 0 Other;
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Pred. No. 1.3
0; Mismatches
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                                                                                                                                                               vaccination; mucokinetic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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RESULT 4
ABK56534/c
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      fibrosis, obstructive bowel syndrome and any other diseases or conditions that are related to or will respond to the levels of CLCA1 in a cell or tissue. The sequences are useful for reducing CLCA1 activity in a cell, hence, are useful for treatment of a patient having a condition associated with the level of CLCA1, where the invention further comprises the use of one or more therapies under conditions suitable for the treatment, for example, oxygen therapy, bronchodilators, corticosteroids, antibacterials, vaccinations, acetylcysteine and mucokinetic agents. The nucleic acids of the invention are also used as diagnostic tools to examine genetic drift and mutations within diseased cells or to detect the presence of CLCA1 RNA in a cell. This sequence represents an enzymatic nucleic acid molecule of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to enzymatic nucleic acid molecules that down regulate expression of chloride channel calcium activated 1 (CLCA1) genes by cleaving RNA derived from the genes. The nucleic acid sequences are useful as pharmaceutical agents for treating conditions such as chronic obstructive pulmonary disease (COPD), chronic bronchitis, asthma, cystic obstructive pulmonary disease (COPD), chronic bronchitis, asthma, cystic obstructive pulmonary disease (COPD).
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                                                                                                                                                                                                    antiinflammatory; chronic obstructive pulmonary disease; COPD; asthma; chronic bronchitis; cystic fibrosis; obstructive bowel syndrome; oxygen therapy; bronchodilator; corticosteroid; vaccination; mucokinet acetylcysteine.
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                                  09-AUG-2000;
                                                                    09-AUG-2001; 2001WO-US024970
                                                                                                      14-FEB-2002
                                                                                                                                      WO200211674-A2
                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                          Human; chloride channel calcium activated 1; CLCA1; ss; antiasthmatic;
                                                                                                                                                                                                                                                                                                            Human CLCA1 gene enzymatic nucleic acid #905.
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                                                                                                                                                                                                                                                                                                                                                                                ABK56534;
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                                  2000US-0224383P
                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                 RNA; 17
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Pred. No.
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thes 0;
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(RIBO-) RIBOZYME PHARM INC.

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RESULT 5
ABK56532/c
ID ABK565
XX ABK565
XX ABK565
XX ABK565
XX Human
XX Human
XX Human
XX Human
XX ACTION
XW ALTION
XW Chroni
XW Chroni
XW Chroni
XW Oxygen
XW ACTION
XW OX900
XW OX900
XX HOMO 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 15
                                                                                                                                                                                                                                                                                    antiinflammatory; chronic obstructive pulmonary disease; COPD; asthma; chronic bronchitis; cystic fibrosis; obstructive bowel syndrome; oxygen therapy; bronchodilator; corticosteroid; vaccination; mucokinetic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Enzymatic polynucleotide that down regulates expression of chloride channel calcium activated gene, useful for treating Chronic obstructive pulmonary disease (COPD), chronic bronchitis and asthma.
                                                                                                                              09-AUG-2001; 2001WO-US024970
                                                                                                                                                                  14-FEB-2002.
                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                       Human;
                                                                                                                                                                                                                                                                                                                                                                                      Human CLCA1
                                                                                                                                                                                                                                                                                                                                                                                                                            02-JUL-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17
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                                                                                         09-AUG-2000; 2000US-0224383P
                                                                                                                                                                                                      WO200211674-A2
                                                                                                                                                                                                                                                                             acetylcysteine.
                                  (RIBO-) RIBOZYME PHARM INC. (SYNT) SYNTEX USA LLC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98
                                                                                                                                                                                                                                                                                                                                chloride channel calcium activated 1; CLCA1; ss; antiasthmatic; flammatory; chronic obstructive pulmonary disease; COPD; asthma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     l Similarity
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                                                                                                                                                                                                                                                                                                                                                                                      gene enzymatic nucleic acid #903.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 15;
Pred. No.
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Best Local S
Matches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     obstructive pulmonary disease (COPD), chronic bronchitis, asthma, cystic fibrosis, obstructive bowel syndrome and any other diseases or conditions that are related to or will respond to the levels of CLCA1 in a cell or tissue. The sequences are useful for reducing CLCA1 activity in a cell, hence, are useful for treatment of a patient having a condition associated with the level of CLCA1, where the invention further comprises the use of one or more therapies under conditions suitable for the treatment, for example, oxygen therapy, bronchodilators, corticosteroids, antibacterials, vaccinations, acetylcysteine and mucokinetic agents. The mucleic acids of the invention are also used as diagnostic tools to examine genetic drift and mutations within diseased cells or to detect the presence of CLCA1 RNA in a cell. This sequence represents an enzymatic nucleic acid molecule of the invention
                                                                                                                                                                                                                                                                                                                                                    Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; murine; tumour suppression; tumour reversion; apoptosis; virus resistance; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to enzymatic nucleic acid molecules that down regulate expression of chloride channel calcium activated 1 (CLCA1) ger by cleaving RNA derived from the genes. The nucleic acid sequences are useful as pharmaceutical agents for treating conditions such as chronic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Enzymatic polynucleotide that down regulates expression of chloride channel calcium activated gene, useful for treating Chronic obstruction pulmonary disease (COPD), chronic bronchitis and asthma.
   New isolated nucleic with tumors and cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Thompson J, Grupe A;
                                                      WPI; 2003-333167/31.
                                                                                                                                                                                                                                  27-MAR-2003
                                                                                                                                                                                                                                                                                                     Mus musculus.
                                                                                                                                                                                                                                                                                                                                     schizophrenia;
                                                                                                                                                                                                                                                                                                                                                                                                                        Murine oligonucleotide associated with tumour supression,
                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-JUL-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ACC68011 standard; DNA; 17
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                                                                                     Telerman A,
                                                                                                                                                         17-SEP-2001; 2001FR-00011979
                                                                                                                                                                                            17-SEP-2002; 2002WO-IB004210.
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                                                                                                                         (MOLE-) MOLECULAR ENGINES LAB
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                                                                                     Amson R,
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                                                                                                                                                                                                                                                                                                                                     88.
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100.0%; Pr
acid, useful for treating viral diseases associated degeneration, also related polypeptides, antibodies
                                                                                      Tuijnder M;
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Disclosure; Page 645; 738pp;

French.

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cells

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RESULT 7
ADC59459
ID ADC59459
ID ADC59459
ID ADC59459
AC ADC5
AC ADC5
XX ADC5
XX Huma
XX Huma
XX Peng
RW Prim
XX JP2(
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PF A (KAK
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Best Local S
Matches 14
               The invention relates to a 450 kD human epidermal autoantigen, epiplakin (ADC59448), and nculeic acids encoding it (ADC59447). The invention also encompasses an epiplakin antigenic epitope (ADC59449) which is reactive with serum from patients with autoimmune disease, fusion polypeptides containing epiplakin or its epitope, an antibody against epiplakin, host cells compresing human epiplakin nucleic acids transgenic animals which under- or over-express epiplakin, epiplakin nucleic acid probes for diagnosis of autoimune disease, and methods of screening for agents which modulate epiplakin activity. Epiplakin polypeptides and polymucleotides are useful in drug screening for agents which promote or inhibit the activity of epiplakin which can be used in the treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to murine oligonucleotides (ACC62754-ACC68806), which are associated with tumour suppression, tumour reversion, apoptosis and virus resistance. The oligonucleotides are useful as (1) as probes and primers for detecting, identifying, quantifying and/or amplifying nucleic acid, e.g. as one component of a gene chip; in vitro as (ant.) sense reagents; and (2) for production of recombinant polypeptides. The oligonucleotides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia
                                                                                                                                                                                                                                                                       Novel protein having epiplakin activity, useful for screening as which inhibit and activate epiplakin activity, and for treating autoimmune skin disease such as pemphigoid or pemphigus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; epiplakin; epidermal autoantigen; autoimmune disease; skin disease; transgenic animals; diagnosis; drug screening; pemphigoid; pemphigoid; pemphigus; dermatological; immunosuppressive; expression analysis; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                pemphigus; dermatological; primer; ss.
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                                                                                                                                                                                                                                            Example 4;
                                                                                                                                                                                                                                                                                                                                                                                                                             16-JUL-2001; 2001JP-00216025
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 disease, particularly those
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                                                                                                                                                                                                                                          ID NO 13; 55pp;
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                                                                                                                                                                                                                                                                                                                                                                                         SHINKO JIGYODAN
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   of the skin
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   such
 as pemphigoid
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RESULT 8
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Matches 14
                                 This oligonucleotide primer was designed to amplify a conserved region of the pleckstrin homology domain of clone 145/H. It was used with a reverse primer (see AAV6425) in the PCR amplification of the CDNA derived from mouse embryo (11 day post-coitus) liver. A PCR product, termed CH7, was subsequently used as probe to screen the embryonic liver library. 2 Overlapping clones were obtained and used to produce a nucleotide sequence (see AAV6414) coding for a novel protein, termed praja-1 (see AAW81641), of the mouse early developing liver. These can be used in the treatment and diagnosis of liver disease and other disorders, including those relating to oncogenesis and tissue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           pemphigus: Epiplakin antibodies and nucleic acid probes are useful diagnosis of these diseases. Sequences ADC59459-ADC59460 represent epiplakin PCR primers used to generate a probe used in expression analysis in an example of the invention.
Sequence
                                                                                                                                                                                                                                         New isolated early liver development genes - used to develop products treating, e.g. liver disease, hepatocellular carcinoma, degenerative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hepatocyte; liver disease; anhidrotic ectoderm dysplasia; hepatocellular carcinoma; colon cancer; spinocerebellar ataxia; sideroblastic anaemia; haemochromatosis; Ito cell; fibrosis;
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                                                                                                                                                                                                 Example 1; Page 28; 92pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Praja-1; mouse; liver development; haematopoiesis; iron transport;
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Query Match Best Local

Similarity

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Score Pred.

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DB 2; 4e+03;

Length

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RESULT 9
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                                                                                                                                                                       isolated during early liver development that are useful in the diagnosis CC and treatment of end stage liver disease and other disorders. Specifically, it refers to genes that encode proteins such as elf 1-3, CC livor-1 (145), pk, protein 106 and praja-1. The present invention CC describes the characterisation of these early liver development proteins, cc and also methods to raise peptide specific antibodies that are useful as markers, as well as for tracing hepatocyte lineage. Furthermore, elf CC proteins 1-3 are useful in treating disorders including cholestasis, cc biliary stones, hepatic obstruction, stricture, primary biliary cirrhosis cr primary sclerosing cholangitis. In addition, through using gene cetoderm dysplasia, hepatocellular carcinoma and other diseases including anaemia, ataxia, degenerative neurological disorders and hemochromatosis. Accordingly, these proteins can be described as chepatotropic, cytostatic, neuroprotective, antianaemic and cardiant. This cligonucleotide sequence is the forward PCR primer used to amplify murine colone 145/PH of the invention.
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Best Local
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                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention relates to novel genes and encoded proteins thereof.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antibodies recognizing early liver development proteins, useful as markers, in identifying peptides and proteins having early liver development characteristics, tracing hepatocyte lineage or treating liver
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30-APR-1998;
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98WO-US008656.
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RESULT 11 AAQ95923/c ID AAQ959

AAQ95923 standard; DNA; 16 BP

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RESULT 10
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Matches 14
                                                                                                                                                                                                                                  (I). (I) is useful as a medicament and in the manufacture of a medicament for treating plakoglobin related diseases, such as skin carcinoma or an acantholytic disease, and to screen composed that interfere with the interaction of the polypeptide with plakoglobin The plakoglobin related diseases include basal cell carcinoma, squamous cell carcinoma, extramammary paget's disease, Naxos disease, heart diseases, skin
                                                                                                                          Sequence
                                                                                                                                                                                     blistering and acantholytic diseases such as subcorneal acanth
Grover's disease, Hailey-Hailey's disease or Darier's disease,
ectodermal dysplasia/skin fragility syndrome. AAS98201- AAS982
                                                                                                                                                                                                                                                                                                                                                                                               New polypeptide, useful for treating skin carcinoma or acantholytic disease such as Grover's and Darier's disease, comprises a protein interacting with human plakoglobin and involved in transduction of plakoglobin related signal to nucleus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; plakoglobin; cytostatic; osteopathic; dermatological; cardiant; plakoglobin related disease; skin carcinoma; acantholytic disease; basal cell carcinoma; squamous cell carcinoma; Naxos disease; PCR primer; extramammary Paget's disease; heart disease; skin blistering; subcorneal acantholysis; Grover's disease; Hailey-Hailey's disease;
                                                                                                                                                                                                                                                                                                                The invention relates to an isolated plakoglobin interacting polypeptide (I). (I) is useful as a medicament and in the manufacture of a medicament
                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 28; 98pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-062246/08
                                                                                                                                                        represent novel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-MAY-2000; 2000EP-00201668
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-MAY-2001; 2001WO-EP004872
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAS98254;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAS98254 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Van Roy F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (VLAA-) VLAAMS INTERUNIVERSITAIR INST BIOTECHNOG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Darier's disease; ectodermal dysplasia; skin fragility syndrome; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human plakoglobin interacting protein PLA_2H12 PCR primer #50.
19
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                                                             14;
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                                                                             Similarity
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                               TCATCTGCCAGGAC
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                                                             Conservative
                                                                                                                          BP; 4 A; 4 C; 12 G; 6 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bonne S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first
                                                                                                                                                         human plakoglobin interacting protein coding of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              entry)
                                                                            100.0%;
                                                                                          10.6%;
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                                                           0;
                                                                             Score 14;
Pred. No
                                                             Mismatches
                                                                             No.
                                                                                                                          U; 0 Other;
                                                                                          DB
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                                                                                          Length
                                                             Indels
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ABK55934/c
ID ABK559
XX ABK559
AC ABK559
XX O2-JUL
XX U2-JUL
XX U2-JUL
XX Human
XX Human
XX Human
KW Antiin
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                          polymorphic, simple sequence repeats and can be used in automated genotyping. esp. fluorescence-based. The primers correspond to the unique DNA sequence surrounding each marker, and PCR is used to detect each polymorphism. When the MMS show considerable polymorphism (ie. a difference in the number of repeats) between individuals, the markers can be particularly informative. The MMS can be ideal for linkage studies. Kits comprise at least 4 groups, of a lost 3 sets, each comprising labelled primers for PCR amplification of the DNA. Group 13 primer pairs are shown in AAQ95915-46. The published size range of the D16S413 allele is 13-149 bp, and the degree of heterozygosity in the population is
Human; chloride channel calcium activated 1; CLCA1; ss; antiasthmatic; antiinflammatory; chronic obstructive pulmonary disease; COPD; asthma; chronic bronchitis; cystic fibrosis; obstructive bowel syndrome;
                                                                                                                                                                                                                                                                                                                                                                    Sequence 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     microsatellite marker sequences (MMS) at approx. 10-50 cM into throughout the human genome which can be detectably labelled.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The method aims to provide a collection of highly reproducible microsatellite marker sequences (MMS) at approx. 10-50 cM intervals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kit for automated genotyping contg. pairs of PCR primers - designed amplify polymorphic nucleotide repeat sequences, arranged in sets exwith a characteristic fluorescence label, useful e.g. in detection of
                                                                                                   02-JUL-2002
                                                                                                                                                              ABK55934 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-DEC-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer; polymerase chain reaction; PCR; linkage study; locus;
microsatellite marker sequence; automated genotyping; allele;
polymorphism; detection; Homo sapiens; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Primer A (Group 13,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21-FEB-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         VINU (OCYU)
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                                                                      CLCA1
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                                                                                                                                                                                                                                                                        79
                                                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                                                                                                                                                        TACTCGGGCTGGA
                                                                                                                                                                                                                                                                                                                                                                    BP; 5
                                                                     gene enzymatic nucleic acid #305.
                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         JOHNS HOPKINS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7M-2; 104pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               genetic rearrangement
                                                                                                                                                                                                                                                                                                                                                                  A
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                                                                                                                                                                                                                                                                                                                                                                  6 C; 3 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                     100.0%;
                                                                                                                                                                                                                                                                                                                                    9.8%;
                                                                                                                                                                                                                                                                        91
                                                                                                                                                                ₽₽
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                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                     Score 13;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  English.
                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                    DB 2;
                                                                                                                                                                                                                                                                                                                     1.3e+04;
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                                                                                                                                                                                                                                                                                                                                 Length 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          arranged in sets each e.g. in detection of
                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                    0
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ABZ65159
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Enzymatic polynucleotide that down regulates expression of channel calcium activated gene, useful for treating Chronic pulmonary disease (COPD), chronic bronchitis and asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
Homo sapiens.
                                                  anti-rheumatic;
                                                                        Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
                                                                                                                                                         Human HER2 DNAzyme substrate #616
                                                                                                                                                                                                              21-MAR-2003
                                                                                                                                                                                                                                                                    ABZ65159
                                                                                                                                                                                                                                                                                                                         ABZ65159 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-AUG-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    oxygen therapy; bronchodilator; corticosteroid; vaccination; mucokinetic; acetylcysteine.
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SYNTEX USA LLC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nucleic acid molecule of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 3 A; 7 C; 3 G; 0 T; 4 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2000US-0224383P
                                                                                                                                                                                                              (first
                                                     cancer;
                                                                                                                                                                                                                                                                                                                      RNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; --
                                                                                                                                                                                                              entry)
                                                                                                                                                                                                                                                                                                                         17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 13;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.3e+04;
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onic obstructive
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RESULT 14
AAH27321/c
ID AAH273
XX AAH273
AC AAH273
XX O8-AUG
DT 08-AUG
XX Tumour
KW Tumour
KW Inflam
KW Cytoki
KW autoim
KW cytoki
KW autoim
KW Cytoki
XX Homo s
XX Homo s
XX OS HOMO s
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, A-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ5989 - ABZ65216, ABZ64544 - ABZ65531. ABZ66520 - ABZ66585 represent substrate/target sequences for the human ribozymes of the invention
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06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
                                                                                                                                                                                                                                                                                                                                                               Tumour suppressor gene 16; TSG16; human; immune response modulator; inflammatory response modulator; signal transduction activator; cytokine inhibitor; gene therapy; anticancer; anti-inflammatory; autoimmune disorder; infection; chromosome 16q24.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   treating cancer, mu
HER2, K-Ras, H-Ras,
                                                       29-OCT-1999;
                                                                                                              30-OCT-2000; 2000WO-AU001329
                                                                                                                                                                                                                             WO200132861-A1
                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                     cellular proliferation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAH27321;
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  (WOME-) WOMEN'S & CHILDREN'S HOSPITAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CCAGGACCUGCUG 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR
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Pred. No. 1.3e+04;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to human tumour suppressor gene 16 (TSG16; see AH23688). TSG16 was isolated from chromosome 16q24.3. TSG16 suppresses cellular proliferation. TSG16 is useful for treating disorders associated with decreased expression or activity of TSG16, e.g. cancers, (auto)immune disorders, inflammation, complications of wound healing and infortions of the relations of the suppression of the s
The present invention relates to human tumour suppressor gene 16 (TSG16; BAH2368). TSG16 was isolated mid from chromosome 16q24.3. TSG16 suppresses cellular proliferation. TSG16 is useful for treating disorder; suppresses cellular proliferation or activity of TSG16, e.g. cancers, associated with decreased expression or activity of TSG16, e.g. cancers,
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                                                                                                                                                                                                                                                           New nucleic acid representing useful e.g. for diagnosis and
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                                            0;
                                                        Score 13;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 13;
Pred. No.
                                                                                                                                                                                                                                                           the human tumor suppressor gene TSG16 treatment of tumors, inflammatory and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          7 T;
                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          fungi, parasites,
is a PCR primer, w
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ou;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 4; Le
1.3e+04;
                                                                    DB 4;
                                                        1.3e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0 Other;
                                                                                                                                                                                                                                                                                                                     Kochetkova
                                            0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 19
                                                                  Length 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          β, protozoa
which was ι
                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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                                            Gaps
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RESULT 17
ADL78982
ID ADL78
XX RNA 11
XX RNA 11
XX ADL78
XX RNA 11
XX ADL78
XX ADL78
XX RNA 11
XX ADL78
XX ADL7
                     CC downregulate expression of one or more human epidermal growth factor CC receptor (EGFR) genes (including HER1, HER2 and HER4) by RNA CC interference. The siNAs may or may not comprise ribonucleotides and may CC be double or single stranded. They further comprise sense and antisense CC regions, or alternatively are assembled from a sense oligonucleotide and CC an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short CC can contain deoxyribonucleotides, and can be chemically synthesised, CC expressed from a vector or enzymatically synthesised. The invention also CC relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are CC used to modulate expression of EGFR genes in cells, tissue explants or CC organisms (e.g., by ex vivo gene therapy), or in grafts and transplants (for the treatment of a variety of conditions. They may be used for CC treating a wide range of cancers such as breast and ovarian cancer. The SiNAs are also useful for drug screening, diagnosis, therapeutic target CC identification and validation, genetic engineering, pharmacogenomics, Studying gene function, and gene mapping (e.g., of single nucleotide).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-JUN-2002;
03-JUL-2002;
29-AUG-2002;
05-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-SEP-2002;
21-OCT-2002;
15-JAN-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of the epidermal gro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          pharmacogenomics; gene function analysis; gene mapping; cancer; cytostatic; human; oncogene; epidermal growth factor receptor; HER2; EGFR2; neu; erbB2; c-erb-B-2; target sequence; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    short interfering RNA; siRNA; double-stranded RNA; micro-RNA; short hairpin RNA; shRNA; expression modulation; gene therapy drug screening; diagnosis; therapeutic target identification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to short interfering nucleic acids (siNA) which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mcswiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-MAY-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-FEB-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO2003070912-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RNA interference;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human HER2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADL78982
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2003-697612/66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           receptor gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3; SEQ ID NO 147; 171pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RIBOZYME PHARM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ç
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (EGFR2) transcript target sequence/siNA upper
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2002US-0363124P.
2002WO-US016840.
2002WO-US0163552.
2002US-0386782P.
2002US-0393924P.
2002US-0406784P.
2002US-0406778P.
2002US-0409293P.
2002US-0409293P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2002US-0358580P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2002US-00277494
function, and gene mapping . The present sequence rep
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 short
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Beigelman L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 interfering nucleic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fosnaugh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 acid; siNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ζ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the epidermal growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Jamison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         receptor; EGFR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            strand,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ: 147
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RESULT 18
ADL79231/c
ID ADL792
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
The invention relates to short interfering nucleic actus vertically downregulate expression of one or more human epidermal growth factor receptor (EGFR) genes (including HER1, HER2 HER3 and HER4) by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short an antisense oligonucleotide. Specifically, the siNAs include short of th
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-MAY-2002;
06-JUN-2002;
06-JUN-2002;
06-JUL-2002;
03-JUL-2002;
29-AUG-2002;
05-SEP-2002;
                                                                                                                                                                                                                                                                                   New short interfering diagnosis of cancer, o
                                                                                                                                                                                                                     Example 3;
                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-697612/66.
                                                                                                                                                                                                                                                                                                                                                                                       Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; short interfering RNA; siRNA, expression modulation; gene therapy; short hairpin RNA; shRNA, expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-AUG-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                    (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-FEB-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             pharmacogenomics; gene function analysis; gene mapping; cancer; cyrostatic; human; oncogene; epidermal growth factor receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human HER2 (EGFR2) siNA lower strand,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADL79231 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19 BP; 4 A; 7 C; 4 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  human HER2 (EGFR2)-targeted double-stranded siNA, which is identical to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      -MAR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EGFR2; neu; erbB2; c-erb-B-2; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15 CCAGGACCTGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11,
                                                                                                                                                                                                                                                             receptor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              transcript
                                                                                                                                                                                                                   SEQ ID NO 396;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  9.8%;
nilarity 84.6%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; 2002US-0363124P.
2002WS-016840.
; 2002US-0163552.
; 2002US-0386782P.
2002US-0393924P.
2002US-0406784P.
; 2002US-0406784P.
; 2002US-0408378P.
; 2002US-0408378P.
; 2002US-0408378P.
; 2002US-0408378P.
; 2002US-0408378P.
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                                                                                                                                                                                                                                                                                                                                                                                         Pavco
                                                                                                                                                                                                                                                           gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            target
                                                                                                                                                                                                                                                                            g nucleic acid, useful
downregulates expressi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27
                                                                                                                                                                                                                                                                                                                                                                                       Beigelman
                                                                                                                                                                                                                171pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 13; I
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                            , useful e.g. i
                                                                                                                                                                                                                                                                                                                                                                                       Fosnaugh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO:396
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .3e+04;
les 0;
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                                                                                                                                                                                                                                                                                                                                                                                       Jamison
                                                                                                                                                                                                                                                                               treatment :
e epidermal
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                                                                                                                                                                                                                                                                        : and
:1 growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 miRNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EGFR
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                        short
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RESULT 19
AAG33460
ID AAG33460
ID AAG33
XX AAG33
XX AAG33
XX AAG33
XX II-DE
DT 11-DE
DT 11-MA
XX PACK
XX HIV;
KW HIV;
KW PART!
XX USN77
XX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             airpin RNA (shRNA). The siNAs can be unmodified or Green Cally Synthesised, can contain deoxyribonucleotides, and can be chemically Synthesised, cxpressed from a vector or enzymatically Synthesised. The invention also CC relates to kits for the in vitro or in vivo delivery of siNA; conjugates cand/or complexes of siNA; and vectors that express siNA. The siNAs are cused to modulate expression of EGFR genes in cells, tissue explants or corganisms (e.g., by ex vivo gene therapy), or in grafts and transplants corect the treatment of a variety of conditions. They may be used for treating a wide range of cancers such as breast and ovarian cancer. The custing a wide range of cancers such as breast and ovarian cancer. The cidentification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide copolymorphisms). The present sequence represents the lower strand of a HER2 (EGFR2)-targeted double-stranded siNA.
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HDPACKI is identical to HIV with 3 important deletions which eliminate an essential packaging signal, part of the env region and the nef and 3' LTR region which is replaced by the polyA site of the SV40 late genes. This construct when coexpressed with the defective RNAs provide proteins for the assembly of new defective virus particles. Example VIII describes the construction of the packaging provirus DNA construct HDPACKI which requires a number of deletions and the addition of a polyA site. The primers of AAQ33459-66 were used. Primers PACK Primer #1 and #2 were used in PCR with pNI4-3 as template. Primer #1 is a deletion primer which removes part of the essential packaging signal. (Note: Revised entry submitted to correct the patent number format of US Government-owned NTIS applications to prevent clashes with ongoing US granted patent numbers.
                                                                                                                                                                                                                                                                                                                                                                                                                   Example VIII;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Defective interfering HIV particles of the amt. of HIV envelope protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Schubert M, Harmison
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11-MAY-1993
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(first en
                                                                                                                                                                                                                                                                                                                                                                                                             Page 52; 110pp; English.
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n surface of host cell.
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                                                                                                                             chimeric protein and comprises a chimeric gene linked to at least 1 HIV LTR (long terminal repeat) promoter sequence, where the chimeric gene comprises a first sequence encoding the ectodomain portion of the CD4 protein and a second sequence encoding the transmembrane and cytoplasmic portions of an envelope protein. The DNA construct may be formulated in a composition together with a carrier and used to treat a patient infected with a human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WWW.
                                                                                                                                                                                                                                                      DNA constructs for inhibiting infectious virus production - containing chimeric gene encoding CD4 ectodomain and viral envelope protein transmembrane and cytoplasmic domains.
                                                                                                                                                                                                    This sequence represents a PCR primer used in the creation of a of the invention. The construct is a DNA construct that encodes
                                                                                                                                                                                                                                                                                               WPI; 1999-059150/05
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28-AUG-1992;
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                                                                                                                                                                                                                                                                                                                  Banjerjea A,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     construct; CD4 fusion protein; LTR promoter; transmembrane g terminal repeat promoter; ectodomain; cytoplasmic domain; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          further information please visit the Derwent web site at .derwent.com/dwpi/updates/ntis_us.html.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              primer PACK#2 used in creation
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                               GCCCTGCATGCAC 14
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                                                                                                                                                                                                                                                                                                                  Schubert
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RESULT

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24-JAN-2002

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ABA81765/c...

ID ABA81765 standard; DNA; 20
XX ABA81765;

XX ABA81765;

XX ABA81765;

XX ABA81765;

XX Aldehyde-dehydrogenase; en
XW Aldehyde-dehydrodiol dehyd
XX Aldehyde-dehydrodiol dehyd
XX Nocardioides sp. KP7.

XX JP2001245662-A.

XX JP2001
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AAD30197/c
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WO200206523-A2
                                                                                                              genotyping; PCR
                                                                                                                                              Human; single nucleotide polymorphism; SNP; diagnosis; pre-disposition; drug induced livertoxicity; screening; UDP-glucuronosyl transferase; UGT1; hepatotoxic reaction; sequence identification; drug metabolism;
                                                                                                                                                                                                                                                                      Human UGT1 gene polymorphism detecting
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                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                      17-MAY-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         s and proteins involved in the polycyclic aromatic compound.
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                                                                                                                     primer; ss.
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PRESENTATION OF A CONTRACT OF 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               also useful in screening assays, to establish animal, cell and models for drug metabolism and for genotyping individuals. The sequence is a common PCR primer used to detect human UGT1 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Detecting predisposition to hepatotoxic reaction of human being caused administration of a compound, by determining single nucleotide polymorphism in UDP-glucuronosyl transferase gene in sample of human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-JUL-2000; 2000EP-00115353
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                                                                                                                                                                                                                                                                                                                                                                               hypotensive; cardiomyopathy; bronchial asthma; gene therapy; vaccine; human; PCR; primer; ss.
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2001US-0274101P.
2001US-0274194P.
2001US-027428P.
2001US-027432P.
2001US-0274349P.
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2001US-0275578P.
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                                                                                                                                                                                                                                                                                                                                                                                                                    cardiant; antiarteriosclerotic; antiasthmatic; cancer;
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Pred. No.
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(UGT1)
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      Padigaru M,
Zerhusen BD
Patturajan !
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19-MAR-2001;
20-MAR-2001;
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31-MAY-2001;
31-MA
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                                         Example C;
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Spaderna SK,
                                                                                                                                                                                                                      Lepley
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2001US-0278152P.
2001US-0278994P.
2001US-0279934P.
2001US-0279934P.
2001US-027993P.
2001US-0279995P.
2001US-02808022P.
2001US-0280802P.
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2001US-0333184P

2001US-033372P

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2001US-0337450P

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2002US-0345705P

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2001US-0294899P.
2001US-0299027P.
2001US-0299303P.
2001US-0299310P.
2001US-0304354P.
2001US-0309198P.
2001US-0312903P.
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2001US-03187402P.
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2001US-0277791P
2001US-0277833P
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Spytek KA, Sh, Gusev V, Ji M, Gangolli E, Casman SJ, Catterton E, Rieger DK; Shenoy i W, Gorman L, Vernet CAM, (Malyankar UM, Burgess C, L SG. Taupier RJ, Pen nan L, Miller CE, CAM, Guo X, Tche ur UM, Gerlach V, C, Leite M, Zho Tchernev V; ch V, Liu Y, , Zhong H, A Pena Kekuda CEA, Anderson Alsobrook 7 5 5 a b

NOVX polypeptides and a disorder associated cancer, hypertension, polynucleotides, useful for preventing or treating with aberrant NOVX expression or activity e.g., atherosclerosis, cardiomyopathy or bronchial

1103pp; English

This invention describes novel human NOVX polypeptides which have

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RESULT 24
ABX97309
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 08-MAR-2001;
08-MAR-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            human; PCR; primer;
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13; Conservative
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102

GTCCCTGTGTCCT 114

Query Match Best Local S Matches 13

Similarity

9.8%;

Score 13; Pred. No.

Conservative

0;

Mismatches

Indels 20

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Gaps

0

Sequence

20

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e G 7 T;

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U; 0 Other; 멂 1.3e+04; ٠. ص 0; Length

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10-JUL-2001;

31-JUL-2001;

16-AUG-2001;

10-SEP-2001;

11-SEP-2001;

27-SEP-2001;

27-SEP-2001;

18-CCT-2001;

18-CCT-2001;

14-NOV-2001;

15-NOV-2001;

16-NOV-2001;

16
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30-APR-2001
02-MAY-2001
03-MAY-2001
15-MAY-2001
16-MAY-2001
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31-MAY-2001
31-MAY-2001
31-MAY-2001
31-MAY-2001
11-JUN-2001
11-JUN-2001
11-JUN-2001
                                    This invention describes novel human NOVX polypeptides which have cytostatic, cardiant, antiarteriosclerotic, antiasthmatic and hypotensive activity. Pharmaceutical compositions comprising the NOVX proteins or nucleic acid molecules or NOVX antibodies are useful for preventing or treating a disorder associated with aberrant NOVX expression or activity e.g. cancer, hypertension, atherosclerosis, cardiomyopathy or bronchial asthma. The products of the invention can be used for gene therapy or in a vaccine. ABX13460.ABX13462 and ABX97185. ABX97593 represent PCR primers polynucleotides represented in ABX97008-ABX97185 which encode the polypeptides represented in ABU65041-ABU65218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Padigaru
Zerhusen
                                                                                                                                                                                                                                                                                                                                                                                                                 NOVX polypeptides and a disorder associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patturajan M, Gangolli
Fernandes ER, Casman SJ
Spaderna SK, Catterton
                                                                                                                                                                                                                                                                                                                        Example
                                                                                                                                                                                                                                                                                                                                                                                         cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lepley
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                                                                                                                                                                                                                                                                                                                      C; Page 646; 1103pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                         hypertension,
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M, Gangolli B,
R, Casman SJ,
Catterton E,
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2001US-0294889P.
2001US-0294899P.
2001US-0299303P.
2001US-0299310P.
2001US-0309198P.
2001US-0312903P.
2001US-0312903P.
2001US-0312903P.
2001US-0312970P.
2001US-0325430P.
2001US-0325430P.
2001US-0325430P.
2001US-0325430P.
2001US-0335301P.
2001US-033531P.
2001US-033531P.
2001US-033531P.
2001US-0335272P.
2001US-0335274P.
2001US-0335274P.
2001US-0337426P.
2001US-0337426P.
2001US-0337426P.
2001US-0337486P.
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2001US-0287424P.
2001US-0288066P.
2001US-0288342P.
2001US-0288528P.
2001US-0291199P.
2001US-0291199P.
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                                                                                                                                                                                                                                                                                                                                                                                       polynucleotides, useful for preventing or treating with aberrant NOVX expression or activity e.g., atherosclerosis, cardiomyopathy or bronchial
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Ji W, Gorman L, Miller CB
Li E, Vernet CAM, Guo X, To
SJ, Malyankar UM, Gerlach V
Nn E, Burgess C, Leite M, Z
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   , Pena CEA,
r CE, Kekuda
TChernev V;
ch V, Liu Y,
, Zhong H, A
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anorexia;

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RESULT 25
ADH13368
                                                                      RESULT 26
ADN62206
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                                                                                                                                                                                                                                                                        # K K K H
                                                                                                                                                                                                                                                                                                This invention relates to a novel method for the prediction, diagnosis, or prognosis of malignant neoplasia by the detection of at least two markers. The invention may also be useful for the development of cyrostatic compounds through the regulation of the expression of a gene or activity of a protein associated with malignant neoplasia. The method is useful for prediction, diagnosis or prognosis of malignant neoplasia such as breast cancer, ovarian cancer, gastric cancer, colon cancer, cesophageal cancer, mesenchymal cancer, bladder cancer or non-small cell lung cancer. The polymucleotides and polypeptides defined in the specification, antisense polymucleotides targeting the polymucleotides, and compounds identified by the screening methods are useful for preventing or treating malignant neoplasia. The disease treated is preferably breast cancer. The present sequence is that of a PCR primer which was used in the exemplification of the invention.
                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                       Query Match
Best Local
                    ADN62206;
                                                     ADN62206 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Predicting, diagnosing or prognosing malignant neoplasia by detecting least two markers, where the markers are genes from one or more chromosomal regions altered in malignant neoplasia,.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-073279/08.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADH13368
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13-FEB-2003; 2003EP-00003112.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human malignant neoplasia-related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; SEQ ID NO 217; 267pp; English.
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                                                                                                                                                        102 GTCCCTGTGTCCT 114
                                                                                                                                                                                                    1 Similarity
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                                                                                                                                      GTCCCTGTGTCCT
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                                                                                                                                                                                                        Conservative
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                                                     DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
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100.0%;
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                                                     ВP
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                                                                                                                                                                                                                       Score 13;
Pred. No.
                                                                                                                                                                                                        Mismatches
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                                                                                                                                                                                                                                       DB 12;
                                                                                                                                                                                                                       1.3e+04;
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                                                                                                                                                                                                                                     Length 20
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                                                                                                                                                                                                        Gaps
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   13-MAR-2001;
13-MAR-2001;
13-MAR-2001;
14-MAR-2001;
16-MAR-2001;
20-MAR-2001;
20-MAR-2001;
20-MAR-2001;
20-MAR-2001;
20-MAR-2001;
20-MAR-2001;
21-MAR-2001;
22-MAR-2001;
23-MAR-2001;
30-MAR-2001;
30-MAR-2001;
31-MAY-2001;
31-MA
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08-MAR-2001;
08-MAR-2001;
09-MAR-2001;
12-MAR-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-JUL-2004 (first entry)
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2001US-027899P

2001US-027934P

2001US-027934P

2001US-0280802P

2001US-0280802P

2001US-0280802P

2001US-0280802P

2001US-028144P

2001US-028144P

2001US-028144P

2001US-028144P

2001US-028806P

2001US-028806P

2001US-029199P

2001US-029199P

2001US-0294489P

2001US-0294899P

2001US-0294899P

2001US-0294899P

2001US-0294899P

2001US-0294899P

2001US-0394899P

2001US-0394899P

2001US-0394899P

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2001US-0394899P

2001US-0394899P

2001US-0394899P

2001US-0395430P

2001US-0318462P

2001US-0318462P

2001US-0318770P

2001US-0335301P

2001US-0335301P

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2001US-0277239P.
2001US-0277321P.
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2001US-0277338P.
2001US-0277791P.
2001US-0277833P.
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2001US-0276000P.
2001US-0276776P.
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2001US-0278894P
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2001US-0275235P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NOVX; diabetes; obesity; infectious disease; an ed cachexia; cancer; neurodegenerative disorder; ease; Parkinson's disease; immune disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3-0275235P.
3-0275578P.
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14-NOV-2001; 14-NOV-2001; 21-NOV-2001; 03-DEC-2001;

2001US-0333184P.
2001US-0333294P.
2001US-0337426P.
2001US-0337426P.
2001US-0337185P.
2001US-0345705P.

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The invention relates to an isolated polypeptide (designated NOVX, or CO NOVI-NOVI27) comprising a sequence selected from 178 fully defined amino acid sequences (and their mature forms, variants and fragments). Also included are an isolated nucleic acid molecule encoding NOVX, a vector comprising the nucleic acid, a cell comprising the vector, methods for determining the presence or amount of the polypeptide or the nucleic acid molecule in a sample, methods for determining the presence of or predisposition to a disease associated with altered levels of expression concept the above polypeptide or nucleic acid molecule in a first mammalian consultation to a method for identifying an agent that binds to the above polypeptide, a method for identifying a potential therapeutic agent for construction or physiological interactions of the polypeptide, a method of capathology that is related to aberrant conscious in the treatment of a pathology that is related to aberrant set of a pathology associated with the polypeptide, a method of cativity or of latency or predisposition to a pathology associated with the polypeptide and a method for modulating the activity of the polypeptide cited above. The composition and methods conscious as a set of the activity of the polypeptide cited above. The composition and methods conscious as a section of the polypeptide cited above. The composition and methods conscious as a section of the polypeptide cited above. The composition and methods conscious as a section of the polypeptide cited above. The composition and methods conscious as a section of the polypeptide cited above. The composition and methods conscious as a section of the polypeptide cited above. The composition and methods conscious as a section of the polypeptide cited above. The composition and methods conscious as a section of the polypeptide cited above. The composition and methods conscious as a section of the polypeptide cited above. The composition and methods conscious as a section of the polypeptide cited above. Th
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Zerhusen BD, (
Patturajan M,
Fernandes ER,
Anderson DW, (
Alsobrook JP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (VERN/)
(GUON/)
(FERN/)
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                                                                                                                                                                                                                                                                                                                            pnarmacogenomics.
                                                                                                                                                                                                                                                                                                                                          preventing or treating NOV
                                                                                                                                                                                                                                                                                                                                                                              New
                                                                                                                                                                                                                                                                                              Example C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (BURG/)
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ZHONG H.
ALSOBROOK J
LEPLEY D M.
RIEGER D K.
BURGESS C E.
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SPADERNA S I
CATTERTON E
LEITE M W.
ZHONG H.
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GUO X S.
TCHERNEV V T.
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KEKUDA R.
PATTURAJAN I
GANGOLLI E J
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SPYTEK K A.
SHENOY S G.
TAUPIER R J.
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CASMAN S J.
MALYANKAR U M.
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JI W.
GORMAN
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                                                                                                                                                                                                                                                                                              SEQ ID NO
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Gusev VY,
Gangolli
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Spaderna SK,
Lepley DM,
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C A M.
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Ji W, Gorman L, Miller CE, Kekuda R
EA, Vernet CAM, Guo XS, Tchernev VT;
J, Malyankar UM, Gerlach V, Liu Y;
SK, Catterton E, Leite MW, Zhong H;
M, Rieger DK, Burgess CE;
                                                                                                                                                                                                                                                                                            786pp; English
                                                                                                                                                                                                                                                                                                                                             Ħ
                                                                                                                                                                                                                                                                                                                                           chromosome mapping, tissue typing
                                                                                                                                                                                                                                                                                                                                                          molecules useful for diagnosing, disorders, e.g. cancer, diabetes,
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RESULT 27
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20-MAR-2001
21-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       diabetes, obesity, infectious diseases, anorexia, cancer-associated cachexia, cancer, neurodegenerative disorders like Alzheimer's disease parkingon'sidisease, immune disorders, haematopoietic disorders, dyslipidaemias, and other chronic diseases. These may also be used i chromosome mapping, tissue typing, preventive medicine and pharmacogenomics. The polypeptides are also useful as vaccines. The present sequence is an RTQ-PCR (real time quantitative PCR) primer uto assay tissue specific expression of a NOVX mRNA.
                                                                                                                                                                                                                                                                                                                                           07-MAR-2002;
                                                                                                                                                                                                                                                                                                                                                                 04-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                            Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                   real
                                                                                                                                                                                                                                                                                                                                                                                                                                           cancer-associated cachexia; cancer; neurodegenerative Alzheimer's disease; Parkinson's disease; immune disorhaematopoietic disorder; dyslipidaemia; chronic diseas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                      US2004043382-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human NOV24b
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                                                                                                                                                                                                                                                                                                                                                                                                                                 quantitative
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           2001US-0276000P.
2001US-027699AP.
2001US-0277339P.
2001US-0277321P.
2001US-0277321P.
2001US-0277318P.
2001US-0277791P.
2001US-0277838P.
2001US-0278152P.
2001US-0278152P.
2001US-0278999P.
2001US-0278999P.
2001US-0279934P.
2001US-0279934P.
2001US-0279934P.
2001US-0280233P.
2001US-0280822P.
2001US-0280822P.
2001US-0280822P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 2 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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2001US-0274849P
                                                                                                                                                                                                                                                                                                                                            2002US-00092900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RTQ-PCR
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2001US-0274281P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NOVX; diabetes; obesity;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              infectious
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                                                                                                                                                                                                                                                                                                                                                                                                                                            disease; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
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Padigaru M, S
Zerhusen BD,
Patturajan M,
Fernandes ER,
Anderson DW,
Alsobrook JP,
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14-NOV-2001;
14-NOV-2001;
14-NOV-2001;
                                                                                                                                             (LEPL/)
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(ANDE/)
(SPAD/)
(CATT/)
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(LEIT/)
(ZHON/)
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(CASM/)
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(GERL/)
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10-JUL-2001;
31-JUL-2001;
16-AUG-2001;
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19-JUN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                       (GUOX/)
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                                                                                                                                                                                                                                ANDERSON D W.
) SPADERNA S K.
) CATTERTON E.
) LEITE M W.
                                                                                                                                                                                                                                                                                                                                               FERNANDES E R. CASMAN S J. MALYANKAR U M.
                                                                                                                                                                                                                                                                                                                                                                                                     GORMAN L.
MILLER C.
KEKUDA R.
PATTURAJAN M.
GANGOLLI E A.
VERNET C. A M.
GUO X S.
TCHERNEV V T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PADIGARU M.
SPYTEK K A.
SHENOY S G.
TAUPIER R J.
PENA C E A.
                                                                                                                                                             ALSOBROOK J P.
LEPLEY D M.
RIEGER D K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZERHUSEN
GUSEV V Y
JI W.
                                                                                                                                                                                                                      ZHONG H.
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2001US-028864P.
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2001US-0288342P.
2001US-029139P.
2001US-02912409P.
2001US-0294889P.
2001US-0294889P.
2001US-029933P.
2001US-029933P.
2001US-039310P.
2001US-039310P.
2001US-0312903P.
2001US-0312903P.
2001US-0312903P.
2001US-0312903P.
2001US-0312903P.
2001US-0312903P.
2001US-0312903P.
2001US-0312903P.
2001US-033231870P.
2001US-033231870P.
2001US-03323184P.
2001US-0333184P.
2001US-0337185P.
2001US-0337185P.
2001US-0337185P.
2001US-0337185P.
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           Taupier RJ, Pena CEA, Li L

man L, Miller CE, Kekuda R;

: CAM, Guo XS, Tchernev VT;

tr UM, Gerlach V, Liu Y;

con E, Leite MW, Zhong H;

KK, Burgess CE;
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WPI; 2004-481158/46.

Wang

Romeo

Suzuki K;

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RESULT 28
ADQ76915
ID ADQ76
DRX PAX RX FXX PXX OS XXX XX DX XX D
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to an isolated polypeptide (designated NOVX, or CC NOV1-NOV127) comprising a sequence selected from 178 fully defined amino CC acid sequences (and their mature forms, variants and fragments). Also CC included are an isolated nucleic acid molecule encoding NOVX, a vector CC comprising the presence or amount of the polypeptide or the nucleic acid molecule encoding NOVX, a vector CC predisposition to a disease associated with altered levels of expression CC included are amethod for identifying an agent that binds to the above polypeptide or nucleic acid molecule in a first mammalian CC subject, a method for identifying a potential therapeutic agent for CC expression or physiological interactions of the polypeptide, a method of agent pathology that is related to aberrant CC expression or physiological interactions of the polypeptide, a method of cc a pathology associated with the polypeptide and a method for modulating CC an pathology associated with the polypeptide and a method for modulating CC are useful for diagnosing, preventing or treating diseases such as CC diabetes, obseity, infectious diseases, anorexia, cancer-associated CC carexia, cancer, neurodegenerative disorders like Alzheimer's disease or Parkinson's disease, immune disorders, haematopoietic disorders. The Composition and other chronic diseases. These may also be used in CC chromosome mapping, tissue typing, preventive medicine and CC pharmacogenomics. The polypeptides are also useful as vaccines. The CC present sequence is an RTQ-PCR (real time quantitative PCR) primer used CC to assay tissue specific expression of a NOVX mRNA.
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                                                                                                                                                                                                                                                                                                                                                                           CA2450504-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                          Escherichia coli.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Biofilm; infection; CsrC; antibacterial; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Escherichia coli csrC
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                                                                                                                                                                                     20-DEC-2002; 2002US-0434779P
                                                                                                                                                                                                                                             22-DEC-2003; 2003CA-02450504
                                                                                                                                                                                                                                                                                                                   26-MAY-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADQ76915;
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                                                                                                                      (KANE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2004-225693/21.
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                                                                                                                      KANE BIOTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene
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Pred. No.
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. 1.3e+04;
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RESULT 29
AAA29645/c
ID AAA296
XX AAA296
XX II-AUG
XX Half-1
KW Half-1
KW Bacill
KW INBect
XX
OS Bacill
KW INBect
XX
OS Bacill
COTYNE
OS Chimer
XX
OS BACIL
OS COYNE
OS CHIMER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence is that of PCR primer DiSerB for the csrC gene CC ADQ76906 of Escherichia coli strain K-12. DiSerB was used with primer CC DiChecki ADQ76916 in an example from the invention for the construction of a csrC null mutant. CsrC interacts with the RNA-binding protein CsrA, CC and antagonises the regulatory effects of CsrA. The invention relates to the csrC gene and RNA, and methods of using these to modulate biofilm CC formation. A claimed method of altering the metabolism or structural or CC functional properties of a bacterial cell comprises altering the genetic CC expression of csrC or the CsrA binding activity of csrC. A result of CC altered genetic expression of csrC may be a change in glycogen CC biosynthesis or gluconeogenesis. A claimed method of reducing biofilm CC cell. A claimed method of inhibiting motility of biofilm-producing CC cell. A claimed method of inhibiting motility of biofilm-producing CC reducing the symptoms of a bacterial infection by biofilm-producing CC bacteria comprises increasing csrC expression. A claimed method of creducing the symptoms of a bacterial infection by biofilm-producing CC bacteria in a mammalian patient involves administering an antibacterial CC agent and decreasing biofilm formation through modulation of csrC.
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Best Local
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                                             Increasing host range microbial insecticide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bacillus thuringiensis.
Corynebacterium diphtheriae.
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                                                                                                                                                                                                                                                                   28-APR-1988;
30-NOV-1992;
                                                                                                                                                                                                                                                                                                                                       10-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bacillus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hybrid toxin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chimeric.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            insecticide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Half-length hybrid toxin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example;
                                                                                                                                                                                                                     (MYCO)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ynthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ü
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13;
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                                                                                                                                                                                                                     MYCOGEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           thuringiensis HD-73; Diphtheria lde; adapter; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTGCGTGAGTTTG 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23
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                                                                                                                                                                   Schwab GE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 5 A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       construction adapter
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92US-00983344
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                                                                                                                                                                                                                   CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            φ
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                                             fusion
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ng bacteria in a mammalian p
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  7 T; 0 U; 0 Other;
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                                                                      of an
                                             of an insecticidal protein used as pesticidal toxin to a cytotoxic a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   No.
                                                                                                                                                                   Wilcox
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toxin;
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fusion protein; chimeric;
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                                                                                                                                                                 Thompson
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Example 1;

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Best Local
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            an insecticidal protein by delivering an insecticidal protein or protein domain to the gut epithelium of a target insect using a targeting protein capable of binding to the gut epithelium. Insecticidal and targeting protein protein do not both originate from Bacillus thuringiensis (B.t.). The hybrid proteins are used as insecticides. The hybrid proteins have increased toxicity against target pests. The present sequence represents an adapter sequence which is used in the construction of hybrid toxins in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A method has been developed of increasing the host range or toxicity of an insecticidal protein by delivering an insecticidal protein or protein domain to the gut epithelium of a target insect using a targeting protein capable of binding to the gut epithelium. Insecticidal and targeting protein do not both originate from Bacillus thuringiensis (B.t.). The hybrid proteins are used as insecticides. The hybrid proteins have increased toxicity against target pests. The present sequence represents an adapter sequence which is used in the construction of hybrid toxins in an example from the present invention
                                                                                                                                               Example
                                                                                                                                                                    Increasing host range microbial insecticide
                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                               Bacillus thuringiensis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bacillus thu insecticide;
                                                                                                                                                                                                                                Culver P,
                                                                                                                                                                                                                                                                                 28-APR-1988;
30-NOV-1992;
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                                                                                                                                                                                                                                                                                                                                             18-APR-2000.
                                                                                                                                                                                                                                                                                                                                                                     US6051556-A.
                                                                                                                                                                                                                                                                                                                                                                                              Chimeric.
                                                                                                                                                                                                                                                                                                                                                                                                                     Corynebacterium
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        construction adapter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first
                                                                                                                                            13; 27pp;
                                                                                                                                                                                                                                                                                 88US-00187167
92US-00983344
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                                                                                                                                                                                                                                                                                                                                                                                                                   diphtheriae
                                                                                                                                                                                                                                Œ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; 5 C; 8 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               entry
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                                                                                                                                                                                                                                  Edwards
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24
                                                                                                                                                                    fusion of
                                                                                                                                            English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   quarter-length hybrid toxin; pesticide;
-73; Diphtheria toxin; fusion protein; chimeric;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  맑
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Pred. No.
                                                                                                                                                                                                                                ΔŢ,
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                                                                                                                                                                    of an insecticidal f pesticidal toxin (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        #9.
                                                                                                                                                                                                                                Wilcox
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.3e+04;
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                                                                                                                                                                    l protein used
to a cytotoxio
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                                                                                                                                                                      cytotoxic
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id toxins in
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Best Local
The invention relates to a hybrid pesticidal protein toxin comprising a cytotoxic agent and a pest gut epithelial cell recognition portion of a protein, where the cytotoxic agent and the recognition portion are not naturally contiguous. The hybrid toxin is expressed by the gene fragment from B.thuringiensis (B.t) var. kurstaki HD-73 which is transformed into a suitable host for use as an insecticide. Expression of the toxin gene results, directly or indirectly in the intracellular production and maintenance of the pesticide. The microbe hosting the toxin gene can be treated under conditions that prolong the activity of the toxin produced in the cell and the treated cell can then be applied to the environment of target pest(s). The resulting product retains the toxicity of the B.t toxin. The present sequence encodes the amino acid sequence of the
                                                                                                                                                                                                                                                                                                                                                         28-APR-1988;
30-NOV-1992;
10-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Corynebacterium diphtheriae.
Bacillus thuringiensis serov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-NOV-2001
                                                                                                                                                                              Novel hybrid pesticidal toxin useful as insecticide, comprises cytotoxic agent, preferably diphtheria toxin and pest gut epithelial cell recognition portion of protein linked by a peptide linker.
                                                                                                                                                                                                                                                                                                           (WILC/)
(EDWA/)
(SCHW/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hybrid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA encoding
                                                                                                                                                         Example
                                                                                                                                                                                                                                                               Wilcox
                                                                                                                                                                                                                                                                                                                                                                                                                               16-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tybrid toxin;
[nsecticide; o
                                                                                                                                                                                                                                         2001-529105/58
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                                                                                                                                                                                                                                                                                    WILCOX E R. EDWARDS D L. SCHWAB G E. THOMPSON M. CULVER P.
                                                                                                                                                         2; Page
                                                                                                                                                                                                                              AAU07529
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              thuringiensis serovar kurstaki;
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т
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Diphtheria toxin
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                                                                                                                                                                                                                                                                                                                                                        88US-00187167.
92US-00983344.
95US-00438465.
                                                                                                                                                         7; 28pp; English.
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/product= '
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/note= "No
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             toxin; diphtheria toxin; cytostatic;
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                                                                                                                                                                                                                                                               PF,
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                                                                                                                                                                                                                                                               Schwab
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          stop or start codon"
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    B-chain/B.t toxin HD-73
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Ω
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1.3e+04;
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RESULT 32
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                28-APR-1988;
30-NOV-1992;
10-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             junction between diphtheria toxin B chain His 48 in a fusion protein between these two proteins. "quarter-length" hybrid toxin, Diphtheria toxin 450-Glu 613
                                                            Novel hybrid pesticidal toxin useful as insecticide, comprises agent, preferably diphtheria toxin and pest gut epithelial cell recognition portion of protein linked by a peptide linker.
                                                                                                                                                                                                                                                                                            CDS
                                                                                                    WPI;
                                                                                                                                         (WILC/)
(EDWA/)
(SCHW/)
(THOM/)
                                                                                                                                                                                                                                                                                                                                Corynebacterium diphtheriae.
Bacillus thuringiensis serovar
                                                                                                                                                                                                                                                                                                                                                        Hybrid toxin;
Insecticide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 24
                                                                                                                   Wilcox
                                                                                                                                                                                                               26-JAN-2000; 2000US-00491320
                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                              DNA encoding
                                                                                                                                                                                                                                                                                                                                                                                               07-NOV-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                              AAS11899
                                              Example
                                                                                                                                                                                                                                              US2001014469-A1
                                                                                                                                                                                                                                                                                                                   Chimeric.
                                                                                                                                                                                                                              16-AUG-2001
                                                                                                                                  (CULV/)
                                                                                             2001-529105/58.
DB; AAU07527.
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The invention relates to a hybrid pesticidal protein to cytotoxic agent and a pest gut epithelial cell recognit protein, where the cytotoxic agent and the recognition naturally contiguous. The hybrid toxin is expressed by from B.thuringiensis (B.t) var. kurstaki HD-73 which is

dal protein toxin cell recognition

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                                                                                                     orientation. Also described is a vector for cloning a DNA molecule comprising: (a) a TAC promoter sequence; (b) a T7 primer sequence; (c) a transcription terminator sequence; (d) a sequence for an operon with an antibiotic resistance marker (ABR); (e) a repressor sequence coding for an inhibitor that is capable of regulating the operon of (d); and (f) a cloning site comprising at least one restriction enzyme site within the repressor sequence that does not change the amino acid sequence of the inhibitor, where the structural genes of the operon are replaced by at least one antibiotic resistance marker and are inhibitor, and where the repressor sequence is operatively linked to the TAC promoter sequence and the T7 primer is introduced for the purpose of sequenceing forms and the T7 primer is introduced for the purpose of sequencing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        a suitable host for use as an insecticide. Expression of the toxin gene results, directly or indirectly in the intracellular production and maintenance of the pesticide. The microbe hosting the toxin gene can be treated under conditions that prolong the activity of the toxin produced in the cell and the treated cell can then be applied to the environment of target pest(s). The resulting product retains the toxicity of the B.t toxin. The present sequence encodes the amino acid sequence of the junction between diphtheria toxin B chain His 484 and B.t HD-73 Cys 10,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cai
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Modified cloning vector with a system for direct selection of recombinants through activation of the antibiotic resistance for molecular DNA cloning for use in molecular biology.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               comprising a ColEl replication origin, a lacl repressor gene, a lac operon-ampr cassette (lac-ABR operon), or an fl origin, all in the same
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention represents a vector (I) for cloning a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-674951/72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cloning vector; molecular cloning; molecular biology; PCR primer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 2; Page 21; 46pp; English.
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       from a forward orientation. Methods and compositions from the present
invention can be used for molecular cloning and for use in molecular
biology. The present sequence represents a PCR primer which is used in
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                              nucleic acid probes and detecting the hybridisation. The nucleic acid probes are attached to a solid support. The analysis comprises monitoring gene expression levels, identifying ballelic markers or polymorphisms, or family members of a gene and a cross-species comparison. Each of the nucleic acids further comprises a tag sequence. The array of nucleic acid probes is useful in in situ hybridisation, in Southern, Northern or dot-blot hybridisation to identify or detect the sequence or specific mutations of any gene, in mapping the 5' termini of mRNA molecules by primer extensions or in screening cDNA or genomic libraries or subclones for additional subclones containing segments of DNA that have been isolated and previously sequenced. The sequence presented is one of the nucleic acid probes incorporated in the microarray. Note: The sequence data for this patent can also be obtained in electronic format directly
                                                                                                                                                                                                                                                                                        Also disclosed is a method of gene expression analysis. The array is used in monitoring gene expression levels by hybridisation to a DNA library, in analysis of genetic variation or in hybridisation of tag-labelled compounds. The nucleic acid probes are specifically designed for analysis of at least one target sequence. The method of analysis comprises the hybridising at least one or more nucleic acids to at least two or more nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                             perfect match, perfect mismatch, antisense match or antisense mismatch Also disclosed is a method of gene expression analysis. The array is use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New array of nucleic acid probes, useful for Southern, Northern or dot-blot hybridization sequence or specific mutations of any gene.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; SEQ ID NO 42570;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    invention discloses a microarray comprising a plurality of nucleic probes including one of 2,018,500 fully defined sequences, or its
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100.0%; Pred. No. 1.3e+04;
ive 0; Mismatches 0;
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Best Local
                                           part of the second retroviral component encodes a protein that is harmful to the cell; a stable cell line that is capable of producing a retroviral vector particle and/or an NOI, where at least part of the second retroviral vector particle of the NOI is harmful to the cell line; a stable cell line that is capable of producing an HIV vector particle; a method of producing a stable cell line compositing infecting a cell with the retroviral vector or particle; and a method of producing a protein that is harmful to a cell by infecting the cell with the retroviral vector or particle. The retroviral vector is useful in medicine, for producing a protein that is harmful to a cell, and for the manufacture of a composition to deliver an NOI to a patient in need of same. This sequence represents a primer used to isolate DNA encoding HIV gag-pol to test for transfer of the genes to a cell using the retroviral vector of
                                                                                                                                                                                                                                                                                          The invention describes a retroviral vector comprising, and capable of expressing, a nucleotide of interest (NOI), which encodes an RNA or production that is harmful to a cell. Also described are: a retrovirus production system for producing the retroviral vector comprising a nucleic acid sequence encoding for the first retroviral vector and nucleic acid sequence encoding for at least one component of the second retroviral vector; a retroviral vector produced by the production system, a retroviral particle obtained or obtainable from the retroviral vector; a cell infected with the retroviral vector and where the NOI or at least
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New retroviral vector comprising, and capable of expressing, a nucleotide of interest (NOI), which encodes an RNA or protein that is harmful to a cell, useful for producing a protein that is harmful to the cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene therapy; retroviral vector; gene expression; retrovirus production system; HIV vector particle; medicine; gag-pol; gene transfer; PCR; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; SEQ ID NO 1; 109pp; English.
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2002US-0417673P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Takeuchi Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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Matches 13
                                                                                                                                      DNA constructs in which the HIV gag, tat and rev genes are expressed from a single promoter and which lack the reverse transcriptase, integrase and envelope proteins are useful to generate immunogens. For the assembly of one suitable construct, the ca. 2kb SacI-MscI DNA fragment from pGGS1 (encoding the gag-protease region of the HIV genome) was subcloned into M13mp18 to provide a template for mutagenesis. Three mutations to remove 2 x HindIII and 1 x PstI sites were simultaneously introduced. A clone containing all 3 mutations was used as a template for further mutagenesis to place a stop codon, a BamHI site, codons for Arg and Ser, a second stop codon and PstI site at the end of the protease gene. Then, the 26-mer AAQ70579 was used to repair a single base deletion and the 39-mer AAQ70580 deleted 19 bases in the 5'-untranslated leader sequence involved in packaging RNA into the virion. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                  DNA construct expresses retroviral GAG-deriv. and REV-like elemetns from same promoter - also the expressed proteins and antigenic particles, useful in vaccines, diagnostic reagnate game the constructions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human Immunodeficiency virus; HIV genome; gag protein; replication-incompetent retrovirus; vaccine; immunogenic particle; ss.
                                                                                                                                                                                                                                                                                                                                                      Example 1; Page 21; 41pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Czaplewski
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12-MAY-1995
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                                                                     Similarity
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                            GCCCTGCATGCAC
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ilarity 100.0%;
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1.3e+04;
                                                                     1.3e+04
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New lentiviral vector and related cells - useful in gene therapy, cacompetent virus.
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                 Human UGT1 gene region 560-587
                                                        30-JUL-2002
                                                                                                                                 ABK51801 standard;
                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes a lentiviral vector (A) comprising
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      quantitative determination of enzymes participating in glucuronic acid conjugation. The method involves the use of oligonuclectide probes hybridising to regions of the human UDF-glucuronosyltransferase (UGT) genes (e.g. UGT1, UGT1A7, UGT1A9, UGT1A10, UGT2A1, UGT2B10, UGT2B11, UGT2B15, UGT2B17, UGT8), and the DDOST gene. The method and probes are useful for the genetic determination of enzymes participating in glucuronic acid conjugation with catalysed UGT. The method is both rapid and accurate. ABK51801-ABK51812 represent regions of the human UGT
Merberg D, I
Spaulding V,
                                                                                                                                                                   26-FEB-1998.
                                                                                                                                                                                                                                               Homo
                                                                                                                                                                                                                                                                                                                                                                             25-MAR-2003
03-AUG-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                     AAV21245;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        or DDOST genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-378271/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; enzyme classification; enzyme quantitative determination; glucuronic acid conjugation; UDP-glucuronosyltransferase; UGT1; ds.
                                                                                           23-AUG-1996;
                                                                                                                                22-AUG-1997;
                                                                                                                                                                                                          WO9807859-A2
                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                   secreted
                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAV21245 standard; DNA; 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to a method for classification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 4; Page 9; 13pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Determination
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-SEP-2000; 2000JP-00272228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-MAR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (SAKA ) OTSUKA SEIYAKU KOGYO KK
                                                      (GEMY ) GENETICS INST INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    being,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CCCTGTGTCCTAC 116
                                                                                                                                                                                                                                                                                                     protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28 BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CCCTGTGTCCTAC 14
                 мссоу JM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                  clone CC365_40 probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        comprises
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                                                                                                                                                                                                                                                                                                                                                                             (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      of enzymes participating in glucuronic acid conjugation comprises use of oligonucleotide probes.
 Jacobs K;
                                                                                           96US-00702344
                                                                                                                                  97WO-US014874
                                                                                                                                                                                                                                                                                                   human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; 10 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 9.8%;
                     Lavallie
                                                                                                                                                                                                                                                                                                     probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 13;
Pred. No.
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                   Ŗ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0 U;
                   Racie LA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 6; Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.3e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0 Other;
                   Treacy M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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RESULT 40
AAX77337/c
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Best Local
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                                                                                                                                                                                                                                                   Jacobs K, r
                               The invention relates to human secreted proteins (AAY21588-597) encoded by polynucleotides obtained from adult retina, adult brain, adult blood or adult testes CDNA libraries. The nucleic acid sequences encoding the secreted proteins correspond to clones AV421-2, BV278-2, C544-1, CC332-33, CC365-40, CG68-4, D329-1, H698-2 and H963-20, (all clones are deposited as ATCC 98145). The PNs and proteins are predicted to have biological activities which would make them suitable for treating, preventing or ameliorating medical conditions in humans and animals, although no supporting data is given. Suggested activities include nutritional activity, cytokine and cell proliferation/differentiation
           activity, immune stimulating (e.g. as vaccines) or suppressing activity, hematopolesis regulating activity, tissue growth activity, activin/
                                                                                                                                                                                                                              WPI; 1999-357814/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New nucleic acid encoding secreted proteins from human cells - useful e.g. as immuno-modulators, antitumour agents, promoters of tissue growth, haemostatic and thrombolytic agents etc.
                                                                                                                                                                            Disclosure; Page 100;
                                                                                                                                                                                                                                                                                                                    21-NOV-1997;
19-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hematopoiesis regulation; tissue growth; chemotactic; thrombolytianti-inflammatory; tumour invasion suppression; gene therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAX77337
                                                                                                                                                                                                                                                                                                                                                                                    03-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                             WO9926973-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Secreted protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-AUG-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1998-169163/15
                                                                                                                                                                                                                                                                                                                                                           20-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The sequence
(Updated on :
                                                                                                                                                                                                     polynucleotides encoding secreted human proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               proli
                                                                                                                                                                                                                                                                                                                                                                                                                                    sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      secreted protein DNA isolating probe.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard; DNA; 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CAGCCCTGCATGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     feration;
                                                                                                                                                                                                                                                      Mccoy JM,
Treacy M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           e is that of a p
25-MAR-2003 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Page 66;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first
                                                                                                                                                                                                                                                                                                                    97US-00976112
98US-00196027
                                                                                                                                                                                                                                                                                                                                                           98WO-US024944
                                                                                                                                                                                                                                                                                              INST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               retina; brain;
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                                                                                                                                                                                                                                                      Lavallie ER,
Spaulding V;
                                                                                                                                                                                                                                                                                             INC
                                                                                                                                                                            106pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           probe used in the correct PI field.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               blood; testis; nutritional activity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   T; 0 U;
                                                                                                                                                                                                                                                                  Collins-Racie LA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               immune stimulation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.3e+04;
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hemostatic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      thrombolytic;
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                                                                                                                                                                                                                                                                   Evans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   human;
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SS; pr
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PR 19-APR
PR 27-SEP
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Best Local 9
                                                              Jacobs K, Merberg D, Kelleher K;
                                                                                                                                                                                                                                                                                 25-OCT-1996;
30-OCT-1996;
13-JAN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                               26-JUL-1996;
23-AUG-1996;
27-SEP-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       autoimmune disorder; multiple sclerosis; systemic lupus erythematosus; rheumatoid arthritis; allergic reaction; asthma; myeloid cell deficiency; lymphoid cell deficiency; osteoporosis; osteoarthritis; peripheral nervous system disease; peripheral neuropathy; Alzheimer's disease; Parkinson's disease; coagulation disorder; Alzheimer's disease; perkinson's disease; coagulation disorder; inflammatory disease; steemic inflammatory response syndrome; SIRS; ischaemia-reperfusion injury; Crohn's disease; anaphylaxis; hypersensitivity; regeneration; neural cell proliferation; fertility;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          thrombolytic activity, receptor/ligand activity, anti-inflammatory activity, cadherin/tumor invasion suppressor activity and tumor inhibition activity. The PNs are also stated to be useful for gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-APR-1996;
19-APR-1996;
                                                                                                                                                                                                                                                                                                                                                  27-SEP-1996
                                                                                                                                                                                                                                                                                                                                                                         27-SEP-1996;
                                                                                                                                                                                                                                                                                                                                                                                        27-SEP-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ss; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human secreted protein cDNA oligonucleotide probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADC38944 standard;
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                                                                                                                                                      (GEMY)
                                                                                                                                                                                                                       1-1997;
-1997;
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                                                                                  Mccoy JM,
Treacy M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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96US-00721923.
96US-00721926.
96US-00738367.
96US-00739775.
97US-00783395.
97US-00833823.
97US-00833823.
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96US-
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96US-00635311.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          human; secreted protein.
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                                                                                  Lavallie ER,
Bowman MR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13
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                                                                                Collins-Racie LA,
Spaulding V, Carl
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1.3e+04;
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WPI; 2003-657236/62

Spaulding

Carlin-Duckett

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RESULT 42
ADH69868/c
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Consteoporosis or osteoarthritis, peripheral nervous system diseases (e.g. peripheral neuropathy, Allaheimer's diseases, Parkinson's disease), coagulation disorders, inflammatory diseases (e.g. systemic inflammatory coagulation disorders, inflammatory diseases (e.g. systemic inflammatory camphylaxis and hypersensitivity. Proteins are also useful for inducing tumour immunity, for inducing bone, carrilage, tendon, ligament and/or regenerating nerve and brain tissue, for inducing fertility and for regenerating nerve and brain tissue, for inducing fertility and for inhibiting tumour growth. Proteins are also useful as chemokine for useful as inhibitors of receptor/ligand interactions. The present sequence represents a oligonucleotide probe for a cDNA encoding a human of the protein are also useful as chemokine for a coagulation of the protein are also useful as chemokine for a coagulation of proteins are also useful as chemokine for a coagulation of proteins are also useful as chemokine for a coagulation of proteins are also useful as chemokine for a coagulation of proteins are also useful as chemokine for a coagulation of proteins are also useful as chemokine for a coagulation of proteins are also useful as chemokine for a coagulation of proteins are also useful as chemokine for a coagulation of proteins are also useful as chemokine for a coagulation of proteins are also useful as chemokine for a coagulation of prot
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Addison's disease; atrophic gastritis; degenerative nervous system disease; multiple sclerosis; degenerative nervous system disease; type I hypersensitivity disease; type I hypersensitivity disease; type II hypersensitivity; Goodpasture's syndrome; allergy; type II hypersensitivity; leprosy; infectious disease; viral infection; type IV hypersensitivity; leprosy; infectious disease; viral infection; HIV; fungal infection; parasitic infection; schetosome; filaria; bacterial infection; Mycobacterium; neoplastic disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              human; T-cell associated disease; Vbeta; autoimmune disease; degenerative nervous system disease; graft versus host disease; hypersensitivity disease; infectious disease; neoplastic disease
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADH69868 standard; DNA; 15
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BD12716 encoded
19-SEP-1994;
                                                                       05-MAR-1999;
                                                                                                                                                                                                                          US2002150891-A1
                                                                                                                                                                                                                                                                                                         Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                    ymphoproliferative disease; leukaemia; lymphoma; cancer;
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                                                                                                                                                                                                                                                                                                                                                                                 cancer;
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94US-00309335
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Mismatches 0;
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Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sclerosis and Alzheimer's disease. Hypersensitivity diseases include Typ I hypersensitivities such as contact with allergens that lead to allergies. Type II hypersensitivities such as those present in Goodpasture's syndrome and Type IV hypersensitivities such as those manifested in leprosy. Infectious diseases include viral infections caused by viruses such as HIV, fungal infections such as those caused by the yeast genus Candida, parasitic infections such as those caused by schistosomes, filaria and bacterial infections such as those caused by Mycobacterium. Neoplastic diseases include lymphoproliferative diseases such as leukaemias, lymphomas and cancers such as cancer of the brain, breast. The present sequence represents a Vbeta gene intron splice site.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kit for diagnozing and treating T-cell associated diseases e.g. autoimmune, degenerative nervous system and infectious disease, comprises nucleic acid primers specifically priming and allowing amplification of a
                                                                                                                                                                                      Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                                                                                                                                                        Human GDMLP-1
                                                                                                                                                                                                                                                                                                                                                    ABN02189 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         including autoimmune diseases, degenerative nervous system diseases, graft versus host disease, hypersensitivity diseases, infectious diseand neoplastic diseases. Autoimmune diseases include Addison's diseastrophic gastritis. Degenerative nervous system diseases include mul
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a kit for diagnosing and treating T-cell associated diseases which comprises a panel of nucleic acid primers specifically priming and allowing amplification of each Vbeta gene, VbetaRNA or CDNA. The kit is useful for diagnosing organ transplant rejection and diagnosing and treating T-cell associated diseases
                                                                                             06-DEC-2001.
                                                                                                                         WO200192524-A2
                                                                                                                                                         Homo sapiens
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(ROWE/) ROWEN L.
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100.0%; Pred. No.
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26-MAY-2000; 21-SEP-2000; 27-SEP-2000;

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25-MAY-2001; 2001WO-US016981.

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ABN02191/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes a human genome-derived myosin-like CC protein 1 (hdDMLP-1). The protein and polynucleotide sequences of hdDMLP-CC 1 can be used in gene therapy and vaccine production. The hdDMLP-1 CC nucleic acids can be used as probes to detect, characterise and quantify CC hdDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hdDMLP-1 CC protein variants having desired phenotypic improvements, and for expressing the proteins. The hdDMLP-1 proteins or polypeptides may be comed as immunogens to raise antibodies that specifically recognise hdDMLP-1 proteins, as standards in assays used to determine the concentration cand/or amount specifically of hdDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser description ionisation, as that appeared to protein, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hdDMLP-1 may be used for diagnosing a CC disorder associated with the expression of hdDMLP-1, in particular heart cand skeletal muscle disorders. hdDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hdDMLP-1 sequence in the expression of hdDMLP-1.

CC The present sequence represents an oligomer used in the screening of the present invention. N.B. The sequence data for this patent did not form part of the printed capture with the screening observations of the printed of the printed sequence as obtained for expression of format directly from WIPO as the present sequence as obtained in electronic format directly from WIPO as the printed sequence in the screening of the printed sequence as obtained for expression of the printed sequence in the screening the screening of the printed sequence in the screening that the screening the screening the screeni
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30-JAN-2001;
30-JAN-2001;
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              Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
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2001WO-US000667,
2001WO-US000668,
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2001WO-US000670,
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The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present invention. N.B. characteristication, but was obtained in electronic format directly from WIPO at figure in the screening of the specification, but was obtained in electronic format directly from WIPO at figure in the printed directly from WIPO.
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21-SEP-2000;
27-SEP-2000;
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RESULT 45 ABN02194/c ID ABN02194 standard; DNA; 17

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The present invention describes a human genome-derived myosin-like protein 1 (hdpMLP-1). The protein and polynucleotide sequences of hdpMLP-1 can be used in gene therapy and vaccine production. The hdpMLP-1 mucleic acids can be used as probes to detect, characterise and quantify hdpMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hdpMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hdpMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hdpMLP proteins, as standards in assays used to determine the concentration and/or amount specifically of hdpMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hdpMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hdpMLP-1 may be used for diagnosing a disorder associated with the expression of hdpMLP-1, in particular heart can also be accepted to the concentration of the particular heart can be accepted to the concentration of the particular heart can be accepted to the concentration of the particular heart can be accepted to the concentration of the particular heart can be accepted to the concentration of the particular heart can be accepted to the concentration of the particular heart can be accepted to the concentration of the particular heart can be accepted to the concentration of the particular heart the concentration o
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present sequence represents an oligomer used in the screening of the MLP-1 sequence in the exemplification of the present invention. N.B. sequence data for this patent did not form part of the printed cification, but was obtained in electronic format directly from WIPO ftp.wipo.int/pub/published_pct_sequence
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                                                                                  The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the representation substrates, to provide initial substrates for the protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLF-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific blomolecule capture probes for surface-enhanced laser description ionisation, as the protein of the prote
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21-SEP-2000;
27-SEP-2000;
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30-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify
                                                                                                                                                          New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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                                                                                                                       Disclosure;
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                                                                                                                                                                                                                                                                                                     Shannon
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RESULT 48
ABNO2192/c
ID ABNO21
XX ABNO21
XX ABNO21
XX Human
DT 29-MAY
XX Human
XX Human
XX Human
XX Human
XX Human
XX Homo &
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CC hGDMLP-1 nucleic acids in samples, as amplification substrates, to CC provide initial substrates for the recombinant engineering of hGDMLP-1 CC protein variants having desired phenotypic improvements, and for CC expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP -1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production and in vaccines or for replacement therapy. The colymucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart CC and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present invention. N.B. CT he sequence data for this patient did not form part of the printed CC specification, but was obtained in electronic format directly from WIPO at figure in the screening of the printed control of the prin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                               30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
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21-SEP-2000;
27-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; genome-derived myosin-like protein 1; GDMLP-1; muscle; myosin; chromosome 22; gene therapy; vaccine; skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human GDMLP-1 17-mer scanning SEQ ID NO:4 sequence
      WPI; 2002-179446/23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-OCT-2000;
30-JAN-2001;
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2001WO-US000661

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2000US-0236359P.
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                                                           Penn
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                                                              Hanzel
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                                                              Rank
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                                                              DR,
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                                                              Shannon ME;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disease;
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RESULT 49
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Best Local
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Grupe
 Thompson Grupe A;
                                                                                                                                                                                                                                                                                                             Human; chloride channel calcium activated 1; CLCA1; 88; antiasthmatic; antiinflammatory; chronic obstructive pulmonary disease; COPD; asthma; chronic bronchitis; cystic fibrosis; obstructive bowel syndrome; oxygen therapy; bronchodilator; corticosteroid; vaccination; mucokinetic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present invention. N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                                                                                                                     Human CLCA1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                            WO200211674-A2
                                                                                                                                                                                                                                                            Homo sapiens.
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                                                                                                                        09-AUG-2000; 2000US-0224383P.
                                                                                                                                                          09-AUG-2001;
                                                                                                                                                                                            14-FEB-2002.
                                                                                                                                                                                                                                                                                              acetylcysteine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABK56932 standard; RNA; 17
                                                    (RIBO-) RIBOZYME PHARM (SYNT ) SYNTEX USA LLC (THOM/) THOMPSON J.
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                                                                                                                                                          2001WO-US024970.
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                 Mcswiggen J,
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                                                                     LLC.
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%; Pred. No. 4.1
0; Mismatches
                 Mckenzie T,
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. 4.1e+04;
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                 Szymkowski DE;
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RESULT 50
ABKS6535/c
ID ABKS65
XX ABKS65
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XX Human
XX Human
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XX Homo s
XW antiin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to enzymatic nucleic acid molecules that down regulate expression of chloride channel calcium activated 1 (CLCA1) genes by cleaving RNA derived from the genes. The nucleic acid sequences are useful as pharmaceutical agents for treating conditions such as chronic obstructive pulmonary disease (COPD), chronic bronchitis, asthma, cystic fibrosis, obstructive bowel syndrome and any other diseases or conditions that are related to or will respond to the levels of CLCA1 in a cell or issue. The sequences are useful for reducing CLCA1 activity in a cell, hence, are useful for treatment of a patient having a condition associated with the level of CLCA1, where the invention further comprises the use of one or more therapies under conditions suitable for the treatment, for example, oxygen therapy, bronchodilators, corticosteroids, antibacterials, vaccinations, acetylcysteine and mucokinetic agents. The nucleic arids of the invention are also used as disconcist that the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Enzymatic polynucleotide that down regulates expression of chloride channel calcium activated gene, useful for treating Chronic obstructive pulmonary disease (COPD), chronic bronchitis and asthma.
                 WPI; 2002-217145/27
                                                    Thompson J, Grupe A;
                                                                                                                                                                          09-AUG-2000; 2000US-0224383P
                                                                                                                                                                                                           09-AUG-2001; 2001WO-US024970
                                                                                                                                                                                                                                                                               WO200211674-A2
                                                                                                                                                                                                                                                                                                                                                acetylcysteine.
                                                                                                                                                                                                                                                                                                                                                                    oxygen therapy; bronchodilator; corticosteroid; vaccination; mucokinetic;
                                                                                                                                                                                                                                                                                                                                                                                      antiinflammatory; chronic obstructive chronic bronchitis; cystic fibrosis; c
                                                                                                                                                                                                                                                                                                                                                                                                  antiinflammatory;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleic acids of the invention are also used as diagnostic tools to examine genetic drift and mutations within diseased cells or to detect the presence of CLCA1 RNA in a cell. This sequence represents an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 4; Page 86; 152pp; English.
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                                                                                                      (RIBO-) RIBOZYME PHARM INC
(SYNT ) SYNTEX USA LLC.
(THOM/) THOMPSON J.
                                                                                                                                                                                                                                               14-FEB-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                    Human; chloride channel calcium activated 1; CLCA1; ss; antiasthmatic;
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hes 0;
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RESULT 51
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to enzymatic nucleic acid molecules that down cregulate expression of chloride channel calcium activated 1 (CLCA1) genes by cleaving RNA derived from the genes. The nucleic acid sequences are useful as pharmaceutical agents for treating conditions such as chronic obstructive pulmonary disease (COPD), chronic bronchitis, asthma, cystic fibrosis, obstructive bowel syndrome and any other diseases or conditions that are related to or will respond to the levels of CLCA1 in a cell or tissue. The sequences are useful for reducing CLCA1 activity in a cell, hence, are useful for treatment of a patient having a condition cassociated with the level of CLCA1, where the invention further comprises the use of one or more therapies under conditions suitable for the treatment, for example, oxygen therapy, bronchodilators, corticosteroids, antibacterials, vaccinations, are also used as diagnostic tools to examine genetic drift and mutations within diseased cells or to detect the presence of CLCA1 maken in a cell. This sequence represents an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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Best Local
                                                         with tumors and cell d
and transfected cells.
                                                                      New isolated nucleic acid, useful for treating viral diseases with tumors and cell degeneration, also related polypeptides,
                                                                                                                       WPI; 2003-313353/30.
                                                                                                                                                       Telerman
                                                                                                                                                                                                                  17-SEP-2001; 2001FR-00011978
                                                                                                                                                                                                                                                 17-SEP-2002; 2002WO-IB004208
                                                                                                                                                                                                                                                                                27-MAR-2003
                                                                                                                                                                                                                                                                                                                WO2003025175-A2
                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                            schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.
                                                                                                                                                                                                                                                                                                                                                                                                          Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; antisense; sense; tumour; cell degeneration; cancer; Alzheimer'
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tumour suppression related human fukutin oligo SEQ ID No 3368
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABT37731 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Enzymatic polynucleotide that down regulates expression of chloride channel calcium activated gene, useful for treating Chronic obstructive pulmonary disease (COPD), chronic bronchitis and asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                      (MOLE-) MOLECULAR ENGINES LAB
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12; Conserv
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100.0%;
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                                                                      associated antibodies
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Disclosure; Page 427;

720pp;

French.

invention relates

to a novel isolated 17 mer nucleic acid

sequence,

The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule, that modulates

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RESULT 52
ABZ65160
ID ABZ65
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Best Local S
Matches 12
                                                                                 Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
                                                                                                                                                                                                                                                                   29-MAY-2001; 2001US-0294140P.
06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras; enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV; anti-rheumatic; cancer; AIDS; ss.
                                               Claim 4; Page 144; 185pp;
                                                                                                                                                           WPI; 2003-140484/13
                                                                                                                                                                                             Mcswiggen
                                                                                                                                                                                                                                                                                                                                           29-MAY-2002; 2002WO-US016840
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human HER2 DNAzyme substrate #617.
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                                                                                                                                                                                                                                 (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14 GGGCTGGATGGA
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                                               English.
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ACD63405/c
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Best Local S
Matches 10
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08-JUN-2001;
24-OCT-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ5989 - ABZ62216, ABZ64544 - ABZ65531, ABZ66520 - ABZ66524, ABZ66530 - ABZ66558 represent substrate/target sequences for the human ribozymes of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                               RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; zinzyme amberzyme; G-cleaver ribozyme; decoy molecule; agener; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HCV minus strand DNAzyme substrate sequence #1044.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ACD63405 standard;
          Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
                                                                                                                                                                                                    (RIBO-)
                                                                                                                                                                                                                                                                                          26-MAR-2001;
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                                                                                                                                                                                                                                                                                                                                                                     WO200281494-A1
                                                                                                                                                                                                                                                                                                                                                                                            Hepatitis C virus.
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                                                                                                                                                                                         (MACE/)
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                                                  2003-229207/22.
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                                                                                                         DRAPER K.
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10; Conser
                                                                                                                                                               BLATT L.
MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
                                                                                                                                                                                                                RIBOZYME PHARM
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                                                                                                                                                                                                                                                                                                                                                                                                                      antiinflammatory; substrate; ss.
                                                                        Macejak D,
Roberts E;
                                                                                                                                                                                                                                      ; 2001US-00817879.
; 2001US-00877478.
; 2001US-0296876P.
; 2001US-0335059P.
; 2001US-0337055P.
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                                                                                    Mcswiggen
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                                                                                    Morrissey
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, C inozymes, zinzymes, amberzymes, and G-cleaver ribozymes. Also disclosed are nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV CC DNA. The nucleic acids may be used to modulate the expression of HBV CC compounds and/or potential therapies directed against HBV, and compounds compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and disease states related to HBV and HCV infection, replication and gene capression such as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence represents a substrate for one of the HCV DNAzyme or minus strand DNAzyme sequences disclosed in the present
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Matches
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08-JUN-2001;
08-JUN-2001;
24-OCT-2001;
                                                                                                                                                                                                                                                                                                                 amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhos liver failure; hepatocellular carcinoma; hepatotropic; cytostatic;
                                       (RIBO-)
(BLAT/)
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(MCSW/)
                                                                                                                                                                                                                                                    WO200281494-A1
                                                                                                                                                                                                                                                                             Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                           Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inc
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                                                                                                                                                                                            26-MAR-2002; 2002WO-US009187.
                                                                                                                                                                                                                         17-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                   HCV DNAzyme substrate sequence #1234.
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              (MORR/)
(PAVC/)
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MACEJAK D.
MCSWIGGEN
MORRISSEY
PAVCO P.
LEE P.
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                                                                                                                                                                                                                                                                                                          antiinflammatory; substrate; ss.
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2001US-0296876P.
2001US-0335059P.
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RESULT 55
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to nucleic acid molecules which modulate Ct the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, C inozymes, zinzymes, and G-cleaver ribozymes, DNAzymes, C inozymes, zinzymes, and G-cleaver ribozymes. Also disclosed are nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV C DNA. The nucleic acids may be used to modulate the expression of HBV compounds and/or potential therapies directed against HBV, and compounds compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and disease states related to HBV and HCV infection, replication and gene expression such as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence represents a substrate for one of the HCV invention or minus strand DNAzyme sequences disclosed in the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Blatt L,
Draper K,
Preparing an amplified product that is free of amplification primer
                                 WPI; 2003-901051/82
                                                               Bradbury AM,
                                                                                                                         24-APR-2002; 2002US-00132067
                                                                                                                                                       11-JUN-2002; 2002US-00167634
                                                                                                                                                                                       06-NOV-2003
                                                                                                                                                                                                                    US2003207282-A1.
                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                           PCR; ss; forensic; paternity testing; disease diagnosis; library; primer
                                                                                                                                                                                                                                                                                                            Amplification
                                                                                                                                                                                                                                                                                                                                          20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                       ADM11727;
                                                                                                                                                                                                                                                                                                                                                                                                   ADM11727 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Page 256; 387pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-229207/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (DRAP/) DRAPER K. (ROBE/) ROBERTS E.
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                                                                                             SOT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TCTGCCAGGACC 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               UCUGCCAGGACC
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Roberts E;
                                                                                             ALAMOS NAT LAB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 2 A;
                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                               Zeytun A;
                                                                                                                                                                                                                                                                                                            method,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     9.1%;
83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8 C; 4 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                           PCR primer #144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mcswiggen J, Morrissey D,
                                                                                                                                                                                                                                                                                                                                                                                                     ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 12; DB 8;
Pred. No. 4.1e+04;
2; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pavco P,
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CC enzyme recognition site is a palindromic sequence or a non-palindromic CC sequence, or is an interrupted sequence. The target polynucleotide CC sequence comprises an antibody variable region or an antibody cCC hypervariable region. The first and second primers specifically hybridise CC to the region encoding an antibody framework region. In preparing a CC library, the library vehicle comprises a cloning vector, particularly a CC phage display vector, a bacterial vector or a yeast vector. Particularly a CC alternatively, the library vehicle comprises a nucleic acid that encodes CC alternatively, the library vehicle comprises an antibody scaffold or a coat protein. The library vehicle comprises an enzyme, a heterologous protein, a green fluorescent protein (GFP) CC scaffold or a coat protein. The library vehicle comprises an incomprise consprises a ribosome. The step of linking the amplified product comprises of ligating an adaptor oligonucleotide to the amplified product to form a CC linked amplified product, and amplifying the linked ligated product. The method is useful in amplified product. The sequences on the amplified product. The method may be used for detecting CC human pathogens, detecting human genetic polymorphisms, detecting RNA and CC mensics, paternity testing, and disease diagnosis. The present sequence CC represents a PCR primer used to demonstrate the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the amplified product with first and second restriction enzymes to obtain the amplified product that is free of amplification primer sequences. Also included is a method of preparing a library, comprising performing the steps of preparing an amplified product that is free of amplification primer sequences and then linking the amplified product to a library vehicle. In the above methods, at least one restriction enzyme cleaves at 2 sites at a distance from its recognition site. At least one restriction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a method of preparing an amplified product that is free of amplification primer sequences comprising amplifying the target nucleic acid sequence in an amplification reaction, and cleaving
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sequences, useful for detecting pathogens, comprises amplifying a target nucleic acid sequence and cleaving the amplified product with restriction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 4;
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            S
                          Matches
                                      Query Match
                                Local
14
                         12;
            68 GCGCGGCTTGCT 79
                                Similarity
                          Conservative
                                100.0%;
                                       9.1%;
                         0
                                Score 1; Pred.
                         Mismatches
                                No.
                                4.1e+04;
                                       DB 11;
                                     Length 17
                         Indels
                         0
                         Gaps
                         0
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Sequence 17 BP; 4 A; 7 C; 6 G; 0 T; 0 U; 0 Other;

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RESULT 56
ADK94931/c
ID ADK949
JP2003259875-A
                                                                   Synthetic
                                                                               human; single nucleotide polymorphism; SNP; ss; primer
                                                                                            Primer of the invention #651
                                                                                                         06-MAY-2004
                                                                                                                     ADK94931;
                                                                                                                                 ADK94931 standard; DNA; 17
                                                                                                       (first entry)
                                                                                                                                 ВP
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08-MAR-2002; 2002JP-00064373

(KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN

08-MAR-2002; 16-SEP-2003

2002JP-00064373

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RESULT 57
AD18675/c
ID AD186
XX AD18675/c
XX AD186
XX AD186
XX BB;
XX BB
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Best Loc
Matches
                                        The invention relates to an enzymatic nucleic acid molecule which specifically cleaves RNA derived from hepatitis C virus (HCV), in which the binding arms of the enzymatic nucleic acid molecule comprises sequences complementary to any of the defined substrate sequences given in the specification. The nucleic acid molecule may be administered for the treatment of HCV infections, especially in combination with type I interferons. The present sequence represents a HCV DNAzyme substrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to a polynuclectide isolated from gene and is useful for detecting a single nuclectide polymorphic human gene or for diagnosing of disease. The invention enables detection of a single nuclectide polymorphism in a human gene. present sequence represents a primer of the invention.
                                                                                                                                                                                                                                                                      Enzymatic nucleic acid molecules which specifically cleave RNA derived from hepatitis C virus (HCV), useful for the treatment of HCV infections, especially in combination with type I interferon therapy.
                                                                                                                                                                                                                                                                                                                                                                                                               Blatt
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HCV DNAzyme
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nucleotide
                         sequence
                                                                                                                                                                                                                                  Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ROBE/)
(PAVC/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18-DEC-2000; 2000US-00740332.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hepatitis C virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (MCSW/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              (MACE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           enzymatic infection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             polynucleotide useful for PCR amplification along with two ent from another set of sequences, or for detecting single otide polymorphism in human gene.
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12; Conser
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ROBERTS E.
PAVCO P A.
MACEJACK D.
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                                                                                                                                                                                                                             SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2000US-00740332.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           acid; RNA cleavage; hepatitis C virus; HCV;
interferon; DNAzyme.
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Pred. No
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Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                            Macejack
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                                                                     Query Match
Best Local S
Matches 10
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                                                                                                                                             Sequence
                                                                                                                                                                                                 The invention relates to an enzymatic nucleic acid molecule which specifically cleaves RNA derived from hepatitis C virus (HCV), in which the binding arms of the enzymatic nucleic acid molecule comprises sequences complementary to any of the defined substrate sequences given in the specification. The nucleic acid molecule may be administered for the treatment of HCV infections, especially in combination with type I interferons. The present sequence represents a HCV DNAzyme substrate
                                                                                                                                                                                                                                                                                                                                                                                      from hepatitis C virus (HCV), useful for the tree especially in combination with type I interferon
                                                                                                                                                                                                                                                                                                                                                                                                        Enzymatic nucleic acid molecules which specifically cleave RNA derived from hepatitis C virus (HCV), useful for the treatment of HCV infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI;
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                                                                                                                                                                                   sequence
                                                                                                                                                                                                                                                                                                                                                       Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Blatt
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(PAVC/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (BLAT/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         enzymatic nucleic infection; type I
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MACEJACK D.
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UCUGCCAGGACC
                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 1234;
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interferon; DNAzyme.
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4.1e+04
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RESULT 59 AAZ95457/c ID AAZ99

a

AAZ95457

standard;

cDNA;

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RESULT 60
AAF75988/c
ID AAF759
XX AAF759
XX 22-MAY
XX 22-MAY
XX Human
XX Human;
KW Cytost
XX Cytost
XX Homo 6
XX Homo 6
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Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes a transcription factor regulating the expression of ethylene-inducible genes in plants, having DNA binding activity specific to the consensus sequence A(T/C)G(A/T)A(C/T)CT. The present invention describes the tobacco ethylene insensitive 3 (EIN3)-like protein, designated TEIL, isolated from Nicotiana tabacum cv Samsun NN. The transcription factor is used to impart environmental stress resistance to plants by transformation with the gene for the transcription factor; and screening potential inhibitors of the expression of ethylene-inducible genes in plants. AAZ95383 to AAZ95476 represent oligonucleotides used in the exemplification of the present
                                                    Human; frizzled cytostatic; PCR
                                                                                               Human
                                                                                                                          22-MAY-2001
                                                                                                                                                                               AAF75988 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Transcription factor regulating the expression of ethylene-inducible genes and gene encoding it, useful for imparting resistance to environmental stress to plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-AUG-1998;
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  WO200112808-A1
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                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 3; Fig 5; 65pp; Japanese.
                                                                                              frazzled family gene 584 PCR primer, SEQ ID NO:18
                                                                                                                                                                                                                                                  14
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12; Conserv
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                                                      family gene 584; embryo; foetus; cancer; drug discovery; primer; ss.
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PS XXX PF XXX PF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity Matches 12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a novel frizzled family gene, 584, from mouse and human (cDNAs given in AAF75973 and AAF75974), and to the mouse and human 584 proteins (AAB73307, AAB73308). Gene 584 is strongly expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel frizzled family genes 584 strongly expressed in embryo and well as in cancer cells, useful in drug development for diseases
                                                                              Novel polynucleotide useful for PCR amplification along with two fragment from another set of sequences, or for detecting single
                                                                                                                                                                                                                                                                                                   08-MAR-2002; 2002JP-00064373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       human; single nucleotide polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Primer of the invention #3055.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              06-MAY-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 abnormal expression including tumor.
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                                                        nucleotide
                                                                                                                                                                               WPI; 2004-093977/10.
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                                                                                                                                                                                                                                        (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN
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Pred. No.
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                                                        gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SNP; ss; primer
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Claim 2; SEQ ID NO 6364; 2627pp; Japanese.

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RESULT 62
ADL95352/C
ID ADL95
XX ADL95
AC ADL95
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XX ADL95
XX PART I
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OS RATT
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Best Local S
Matches 12
                                                                                                                                Treating or ameliorating neurological disorders e.g., cancer pain, osteoarthritis pain, allodynia or hyperalgesia comprises injecting into subject a double-stranded RNA that inhibits the expression of a target
                                                                                                                                                                                                                                                                                                                                                                                 04-SEP-2002;
27-MAR-2003;
    The present invention de neurological disorders,
                                                                                                                                                                                                                              WPI; 2004-248370/23
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                                                                       Example;
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  invention describes a method for treating of disorders, which comprises injecting into
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2003US-0457971P.
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/note= "phosphorothioate linkages"
10. .18
/*tag= b
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/note= "2'.-O-(2-methoxyethyl) ribonuclectide, where C = 2'.-O-(2-methoxyethyl) 5-methyl cytidine and T = 2'.-O-(2-methoxyethyl) 5-methyl uridine"
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100.0%; Pr
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           Treating or ameliorating neurological disorders e.g., cancer pain, osteoarthritis pain, allodynia or hyperalgesia comprises injecting subject a double-stranded RNA that inhibits the expression of a target subject and the expression of a target subject and the expression of a target subject and target subject subj
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                                                                                                                                               Dorn
                                                                                                                                                                                          SAON)
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27-MAR-2003; 2003US-0457971P
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Synthetic
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12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "2'-O-(2-methoxyethy1) ribonucleotide, 2'-O-(2-methoxyethy1) 5-methy1 cytidine and T methoxyethy1) 5-methy1 uridine"
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/note= "2'.-O-(2-methoxyethy1) ribonucleotide,
2'.-O-(2-methoxyethy1) 5-methy1 cytidine and T
methoxyethy1) 5-methy1 uridine"
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PHARMA
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                                                                                                                                                                                          GMBH.
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in; osteoarthritis pain; allodynia; hyperalgesia;
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Example;

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                                                                                                                                                                                                                                                                                   29-JAN-1992;
24-MAR-1992;
12-NOV-1992;
The sequences given in AAQ47539-50 are probes which were used in the quantification of human GTP binding protein (G protein)-specific mRNAs. These probes are based on sequences derived from human and rat G-proteins. These probes were used in the method of the invention for the detection and quantification of mRNAs in a sample without the need to purify the mRNA from cells. The claimed method comprises identifying a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Probe; quantification; human; GTP binding protein; G protein; alpha subunit; specific mRNA; detection; hybridisation; diagnosis; pathophysiology; disease state; hereditary; cancer; infectious;
                                                                                                         Example
                                                                                                                                   Quantitating messenger RNA in sample - using immobilised-polynucleotide having sequence complementary to sequence unique to the MRNA.
                                                                                                                                                                                 WPI; 1993-258695/32.
                                                                                                                                                                                                             Akitaya
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
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26-JAN-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RATBPGTPD/Go-1397 Go specific probe
                                                                                                                                                                                                             'n
                                                                                                                                                                                                                                                      HITACHI
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                                                                                                                                                                                                                                        HITACHI CHEM CO LTD.
HITACHI CHEM RES CEN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GGCTGGATGGAG 96
                                                                                                                                                                                                             Cooper A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 4 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (revised)
(first entry)
                                                                                                                                                                                                                                                                                     92US-00974409
                                                                                                                                                                                                                                                                                                  92US-00827208.
92US-00857059.
                                                                                                                                                                                                                                                                                                                                             93WO-US000977
                                                                                                                                                                                                                                                                                                                                                                                                                                                               pituitary tumour; chain reaction; s
                                                                                                      55; 177pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        9.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                              Mitsuhashi M;
                                                                                                                                                                                                                                           CENT INC
                                                                                                        English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            acromegaly; melanoma cells; diabetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 12;
4.1e+04;
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RESULT 65
AAV17327
ID AAV17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cc polynucleotide sequence unique to the mRNA, and immobilising an oligomer cc complementary to this sequence to an insoluble support. The sample is then incubated with the insoluble support such that the unique sequence cc will hybridise to the bound oligomer and be immobilised. Non-immobilised components are washed from the support and bound RNA is labelled in such ca way that the label is incorporated onto the support relative to the camount of mRNA on the support. The amount of bound label is then cd etermined. This method can be used for the reliable, rapid, simultaneous cquantification of multiple varieties of mRNA. It may be used for cd dagnosing and recognition of pathophysiology of various disease states, eg. hereditary diseases, cancer, and infectious diseases. G proteins are thought to be involved in causing various disease states. A genetic certiciency of Gs protein is the molecular basis of hereditary of cs proteins of proteins are also involved in invasive and meteastatic melanoma cells, and diabetes. See also AAQ47381-666.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                          This sequence is a primer that was used to construct the antibody of the invention. The antibody is an anti-CEA (carcinoembryonic antigen) antibody (806.077 Ab). Host cells or transgenic organisms transformed with DNA encoding the antibody, are used to make the antibody or conjugate. The conjugate is used in a medicament suitable for intravenous administration. The conjugate can be used for cancer therapy, selectively killing tumour cells. The antibody can be used for in vivo or in vitro
                                                                                                                                                                                                                                                                                                                                  04-MAY-1996;
14-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19
                                                                                                                                                          Example
                                                                                                                                                                                                     Anti-carcinoembryonic antigen antibody 806.077
                                                                                                                                                                                                                                                                     Copley
                                                                                                                                                                                                                                                                                                                                                                                 29-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                               13-NOV-1997.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cancer diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Anti-CEA antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Primer used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-JUN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAV17327 standard;
                                                                                                                                                                                                                                      WPI; 1997-558987/51
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                                                                                                                                                                                                                                                                                                    (ZENE ) ZENECA LTD.
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                                                                                                                                                        48; Page 170; 208pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                         cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         construction of
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97GB-00003103.
                                                                                                                                                                                                                                                                                                                                                                                 97WO-GB001165
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        carcinoembryonic antigen; 806.077 Ab; cancer therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antibody
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         of the invention
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cancer

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RESULT 66
RAV41791
ID AAV41791
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     RESULT 67
ADF84430
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Best Local
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Best Local :
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                                                                                                                                                                                                                                                                                                                   The primers AAV41785-V41794 were used in the cloning of human pancreatic carboxypeptidase B (CPB). The co-expression of a modified pro-domain of CPB from a seperate gene enhances enhance expression. This process can be used to produce recombinant CPB in eukaryotic cells, or fusions of CPB with antibody chains. CPB is used in insulin production and protein seguencing, while its fusions with antibody are useful in antibody-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New modified pro-domain of carboxy-peptidase B - enhances expression co-expressed proteins for production of recombinant carboxy-peptidase its fusions with antibodies, used, e.g. in enzyme prodrug therapy.
                                                                                                                                                                                                                Sequence 19
                                                                                                                                                                                                                                                                directed enzyme prodrug therapy. The Modified pro-domain provide increased yields of recombinant CPB, possibly by protecting the C-terminus against enzymatic degradation or by increasing intracellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9835988-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 51; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1998-467168/40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-FEB-1997;
18-OCT-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ss; primer; PCR; amplification; human; pancreatic carboxypeptidase
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ZENE )
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                                                                                                           GGACCTGCTGCA 29
                                                                         GGACCTGCTGCA 12
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                                                                                                                                              Conservative
                                                                                                                                                                                                                  BP; 3
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97GB-00022003.
97GB-00022727.
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                                                                                                                                                                                                                5 C;
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                                                                                                                                            Score 12; DB; Pred. No. 4.1
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Pred. No.
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ADF84749/ID ADF8
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                     Human ABL1-targeted
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Query Match
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Matches S
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29-AUG-2002;
05-SEP-2002;
09-SEP-2002;
14-JAN-2003;
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11-MAR-2002;
06-JUN-2002;
                                                                                                                                                                                                                                                                                                                         nucleic acid (siNA) that downregulates expression of the breakpoint cluster region-v-abl Abelson murine leukaemia viral oncogene homologue 1 (BCR-ABL) gene. The siRNA of the invention demonstrates cytostatic activity and may be useful for modulating expression of the BCR-ABL gene, as well as for treating leukaemia or lymphoma and in diagnosis, drug screening, target identification and validation, genetic engineering, gene function studies and gene mapping. The current sequence is that of the human ABL1-targeted siRNA of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         short interfering nucleic acid; siNA; breakpoint cluster region; v-abl Abelson murine leukaemia viral oncogene homologue 1; BCR-ABL; cytostatic; leukaemia; lymphoma; human; ss; siRNA; ABL1.
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and
                                  ADF84749;
                                                                                                                                                                                                                                                                                                Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a novel double-stranded short interfering
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example
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 26-FEB-2004
                                                                     ADF84749
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             double-stranded interfering nucleic acid, useful e.g. diagnosis of leukemia and lymphoma, downregulates the ster region-Abelson (BCR-ABL) gene.
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                                                                                                                                                                                           59
                                                                                                                                                          7
                                                                                                                                                                                                                            Similarity 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              7; SEQ ID NO 724; 197pp; English.
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                                                                     standard; RNA; 19
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2002US-0409293P.
2003US-0439922P.
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2002US-0386782P.
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entry)
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75.0%;
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Pred. No. 4.1e+04;
3; Mismatches 0
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SiRNA -

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RESULT 69
AAZ05043/c
ID AAZ050
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AC AAZ050
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XX
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Vaccin
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Chlamy
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Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a novel double-stranded short interfering nucleic acid (siNA) that downregulates expression of the breakpoint cluster region-v-abl Abelson murine leukaemia viral oncogene homologue 1 (BCR-ABL) gene. The siRNA of the invention demonstrates cytostatic activity and may be useful for modulating expression of the BCR-ABL gene, as well as for treating leukaemia or lymphoma and in diagnosis, drug screening, target identification and validation, genetic engineering, gene function studies and gene mapping. The current sequence is that of the human ABL1-targeted siRNA of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-SEP-2002;
09-SEP-2002;
14-JAN-2003;
                                                                   paratrachoma; inclusion conjunctiviti
nongonococcal uretritis; epidymitis;
            Chlamydia trachomatis
                                                        bartholinitis;
                                                                                                                                                     07-OCT-1999
                                                                                                                                                                                                           AAZ05043 standard;
                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19
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and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       short interfering nucleic acid; siNA; breakpoint cluster region; v-abl Abelson murine leukaemia viral oncogene homologue 1; BCR-A cytostatic; leukaemia; lymphoma; human; ss; siRNA; ABL1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example
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15-AUG-2002;
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                                                                                                                       primer used to amplify an ORF of Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                double-stranded interfering nucleic acid, useful e.g. diagnosis of leukemia and lymphoma, downregulates the ster region-Abelson (BCR-ABL) gene.
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                                                                                               eye disease;
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2002US-036782P.

2002US-0346782P.

2002US-0404039P.

2002US-0406784P.

2002US-0408378P.

2002US-0409293P.

2002US-0419923P.

2003US-0419923P.

2003US-0419923P.
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                                                      pneumopathy; venereal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NO 1043;
                                                                                                                                                                                                                                                                                                                                              9.1%;
                                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                                                                                                                      3 C; 7 G; 0 T; 3 U; 0 Other;
                                                conventional trachoma; nonendemic trachoma; on conjunctivitis; genital disease; perihepatitis; is; epidymitis; cervicitis; salpingitis; PCR prime pathy; venereal lymphogranulomatosis; ss.
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Pred. No.
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RESULT 70
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17-DEC-1997;
04-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR primers AAZ01426-Z06209 were used to amplify open reading frames (ORFS) of the genome of Chlamydia trachomatis (see AAZ01425). These ORFS encode polypeptides (see AAY3673-437349) which can be used as vaccines against Chlamydia trachomatis. Antisense and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases such as conventional trachoma, nonendemic trachoma, paratrachoma, and inclusion conjunctivitis; genital diseases such as monogenococcal uretritis, perincepathy in breast feeding infants; and venercal lymphogranulomatosis. The polypeptides of the invention may be of use in treating these
                                                                                                                                                                                                                                                                                                                                                                                                                               Respiratory disease; pneumonia; bronchitis; heart disease; sinusitis; purulent otitis media; erythema nodosum; pharyng
                                                                                             21-NOV-1997;
04-NOV-1998;
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                                             (GEST ) GENSET
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97FR-00016034.
98US-0107077P.
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98US-0107078P.
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4.1e+04;
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Griffais R;

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AAX91991-X97517 represent PCR primers used to amplify open reading fram and other nucleic acid sequences from the genome of Chlamydia pneumonia (see AAX91990). C. pneumoniae causes respiratory disease such as pneumonia and bronchitis and is thought to be a contributing factor in heart disease, sarcoidosis, sinusitis, purulent otitis media, crythema nodosum or pharyngitis. The polypeptides encoded by the open reading frames of the C. pneumoniae genome (see AAX34584- AAX35879) can be used
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04-NOV-1998;
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                                                                                                                  Page 1778; Disclosure; 1912pp; English.
                                                                                                                                                                            WPI; 1999-357842/30
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                                                                                                                                              sequence of Chlamydia pneumoniae
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        targeted to the 5'-untranslated region or 3'-untranslated region of a nucleic cranslational termination region or 3'-untranslational start site, it acid encoding a p38 mitogen activated protein kinase (MAPK), where the antisense oligonucleotides inhibit the expression of MAPK), where the antisense oligonucleotides inhibit the expression of MAPK), where the antisense oligonucleotides inhibit the expression of MAPK), where the AAC79480 and AAC79500 and AAC7953 - AAC79570 represent human p38alpha MAPK and p38beta MAPK cDNA sequences. AAC79510 - AAC79500 and AAC7953 - AAC79572 and p38alpha antisense oligonucleotides, while AAC79502 - AAC79521 and antisense oligonucleotides AAC79502 - AAC79523 and antisense oligonucleotides AAC7953 - AAC7953 and antisense oligonucleotides AAC7953 and antisense oligonucleotides are ap8alpha cDNA sequence AAC79523 and antisense oligonucleotides are given in AAC79537 and antisense oligonucleotides have antirheumatic; antiarthritic; immunosuppressive; cardiant and antiinflammatory activity. The antisense oligonucleotides have antirheumatic; antiarthritic; calls or tissues. The oligonucleotides are used for treating an animal with diseases such as inflammatory or autoimmune diseases e.g. rheumatoid arthriting or heart disease. The oligonucleotides are also useful for inhibiting the expression of p38 MAPK in the particle of the p
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                                                                                                                                                                                                                                                                                                                                                                                        This invention relates to antisense compounds 8-30 nucleobases in length
                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 19; Page 63; 90pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antisense oligonucleotide; p38 mitogen activated protein kinase; MAPK; antirheumatic; antiarthritic; immunosuppressive; cardiant; heart disease; antirhelammatory; autoimmune disease; rheumatoid arthritis; apoptosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human p38beta antisense oligonucleotide SEQ
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                                                                                                                                                                       PCR primers AAF80035-36 were used to detect the genotype of human Cf families. The specification describes an odorant binding polypeptide (OBP). OBPs provide long-term retention (gradual release) of lipophilic compounds, so prolong the 'hold' of perfumes, deodorants etc. OBP polypeptides are used as binding proteins for hydrophobic ligands (particularly odorants); as competitive inhibitors (agonists or controlling the controlling proteins for hydrophobic ligands (particularly odorants); as competitive inhibitors (agonists or antipodies for diagnosis of allergy, asthma or cancer; for controlling coldinistion of an odorant, specifically in perfumes, cosmetics or diainfectant compositions, to screen compounds, especially odorants or flavours, e.g. human pheromones, for binding to OBP, also in analysis of complex perfume mixtures; to solubilite lipophilic compounds; for treating hyperlipidemia or obesity, or to supplement non-maternal milk when combined with nutritional fatty acids, as food additives; as a collayed release) but also for delivery across the nlacental barvier (e.g., collayed release) but also for delivery across the nlacental barvier (e.g., collayed release) but also for delivery across the nlacental barvier (e.g., collayed release) but also for delivery across the nlacental barvier (e.g., collayed release) but also for delivery across the nlacental barvier (e.g., collayed release) but also for delivery across the nlacental barvier (e.g., collayed release) but also for delivery across the nlacental barvier (e.g., collayed release) but also for delivery across the nlacental barvier (e.g., collayed release) but also for delivery across the nlacental barvier (e.g., collayed release) but also for delivery across the nlacental barvier (e.g., collayed release) but also for delivery across the nlacental barvier (e.g., collayed release) but also for delivery across the nlacental barvier (e.g., collayed release) but also for delivery across the nlacental barvier (e.g., collayed the delivery across the nlacent
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                                                                                                                    delayed release) but also for delivery across the placental barrier (e.g for detoxification of the foetus); as a marker of pregnancy or foetoplacental pathology (rupture of the amniotic membrane); and as
                                                            Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New human odorant-binding proteins, useful for solubilizing lipophilic compounds in the transportation of anticancer agents or for slow releases
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                                                                                                                                                                                                The specification describes a human embryo-derived (EBD) cell culture that comprises cells which simultaneously express polypeptide or mRNA markers that are characteristic of at least 2 different cell types. The cell types are ectodermal, mesodermal, and/or endodermal cells. The markers include nestin, vimentin, myf6, myosin light-chain 2 ventricular isoform, flk1, alpha-1 fetoprotein and GATA-4. The EBD cells are useful in cell culture, tissue transplantation, tissue engineering, drug discovery and gene therapy. They are especially useful in transplantation therapies for the treatment of Parkinson's disease, amyotrophic lateral sclerosis, stroke, injury to motor neurons, including spinal cord injury, and disbetes. Human EBD cells can be implanted into the central nervous
                                                     system for the treatment of disease or physical brain injury, such as ischemia or chemical injury, as well as to treat diseases of cardiac, skeletal or smooth muscles, liver, pancreas, eyes, vascular diseases, trepopulate bone marrow, in organotypic co-culture, in the biosynthetic production of macromolecules, and in the construction and testing of human artificial chromosomes. AAH42612-85 represent PCR primers and probes for markers of human cells. They were used to examine the mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human embryo-derived cell culture comprising cells that simultaneously express polypeptide or mRNA markers that are characteristic of at least cell types, useful in tissue transplantation, drug discovery, gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               endodermal cell; nestin; vimentin; myf6; flk1; alpha-1 fetoprotein; myosin light-chain 2 ventricular isoform; GATA-4; cell culture; tissue transplantation; tissue engineering; drug discovery; gene therap; transplantation therapy; Parkinson's disease; spinal cord injury; amyotrophic lateral sclerosis; stroke; motor neuron injury; diabetes; brain injury; ischemia; cardiac disease; skeletal disease; eye disease; smooth muscles disease; liver disease; pancreas disease; cell marker;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 7;
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                                      expression
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                                      of human cells
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                                                                                                                                                                         The present invention relates to a new method of interfering with G2/M C checkpoint function in cells exposed to a genotoxic stress thus cinhibiting p53-mediated apoptosis in a cell exposed to genotoxic stress. The method of the invention involves inhibiting p38 mitogen-activated protein kinase (MAPK) activity in the cell and thus interfering with G2/M checkpoint (cell division) function. The invention can be used for interfering with G2/M checkpoint function in cells (e.g. tumour, non cultumour or cardiac cells) exposed to a genotoxic stress, and thus chibiting p53-mediated apoptosis in a cell exposed to genotoxic stress. The method of the invention can also be used for treating a tumour in a culturation can also be used for treating a tumour in a culturation can also be used for treating a condition can concern the cell exposed to genotoxic stress. The methods of the invention are suitable for enhancing the efficacy and safety of genotoxic interapy. The present mucleic acid sequence represents synthetic p38 beta antisense (AS) oligonucleotide concerns the methods of the invention for antisense suppression of p38 MAPK expression in human cancer cells
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Interfering with G2/M checkpoint function in cells exposed to genotoxic stress for treating tumors, by inhibiting p38 mitogen-activated protein kinase activity in cell, thus interfering with G2/M checkpoint function.
                                                                                                                                              Sequence 20 BP; 2 A; 5 C; 8 G; 5 T; 0 U; 0 Other;
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5 US-10-098-26318-42570
0 US-09-866-108-2183
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US-99-866-108-2185
US-99-866-108-2186
US-99-97-046-1303
US-99-97-046-303
US-99-97-046-303
US-99-97-940-332-1234
US-10-669-841-382-3321
US-10-669-841-382-3321
US-10-669-841-382-3321
US-10-723-361-2183
US-10-723-361-2186
US-10-723-361-2183
US-10-723
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Sequence 10682, A
Sequence 4270, Appl
Sequence 2183, Appl
Sequence 2183, Ap
Sequence 2184, Ap
Sequence 2185, Ap
Sequence 2186, Ap
Sequence 2186, Ap
Sequence 2186, Ap
Sequence 2187, Ap
Sequence 1100, Ap
Sequence 1101, Ap
Sequence 2181, App
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15 US-10-153-244-287
15 US-10-120-39-65
15 US-10-271-078-68
15 US-10-350-516-51
15 US-10-350-516-51
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16 US-10-351-846-21
17 US-10-351-846-21
18 US-10-651-722-48
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19 US-09-866-108-5113
9 US-09-866-108-5123
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Sequence 5112,
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ALIGNMENTS

RESULT 1 US-09-927-046-304/c ; Sequence 304, Application US/09927046

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RESULT 3
US-09-927-046-905/c
Sequence 905, Application US/09927046
Publication No. US20030064946A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, In
APPLICANT: McSwiggen, Jim
APPLICANT: Thompson, Jim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Publication No. US20030064946A1
; GRMERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: NcSwiggen, Jim
APPLICANT: Thompson, Jim
APPLICANT: Thompson, Jim
APPLICANT: Ayers, Dave
APPLICANT: Szymkowski, Edmund
TITLE OF INVENTION: Method and Reagent for t
TITLE OF INVENTION: Channel-1
FILE REFERENCE: 249/021
CUCRENT APPLICATION NUMBER: US/09/927,046
CUTRENT FILING DATE: 2001-08-09
NUMBER OF SEQ ID NOS: 5450
SOFTWARE: PATENTIN Version 3.0
SEQ ID NO 304
                                                                                                                                                                                                                                                                                          ; TYPE: RNA; Homo sapiens US-09-927-046-904
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; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-927-046-304
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US-09-927-046-904/c
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CURRENT FILING NUMBER: US/09/927,046
CURRENT FILING DATE: 2001-08-09
NUMBER OF SEQ ID NOS: 5450
SOFTWARE: Patentin version 3.0
SEQ ID NO 904
LENGTH: 17
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Publication No. US20
GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Ayers, Dave
APPLICANT: Grupe, Andrew
APPLICANT: Szymkowski, Edmund
TITLE OF INVENTION: Method and
TITLE OF INVENTION: Channel-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Ribozyme Pharmaceuticals,
APPLICANT: McSwiggen, Jim
APPLICANT: Thompson, Jim
APPLICANT: McKenzie, Tim
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5. US20030064946A1
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Pred. No.
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CURRENT APPLICATION NUMBER: US/09/927,046
CURRENT FILING DATE: 2001-08-09
NUMBER OF SEQ ID NOS: 5450
SOFTWARE: Patentin version 3.0
SEQ ID NO 903
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LENGTH: 17
                                                                                                                                                              Sequence 305, Application US/09927046 Publication No. US20030064946A1
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Best Local
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ORGANISM: Homo sapiens
-09-927-046-905
                                                                                                          APPLICANT: Ribozyme Pharmaceuticals, APPLICANT: McSwiggen, Jim
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APPLICANT: Grupe, Andrew
APPLICANT: Szymkowski, Edmund
TITLE OF INVENTION: Method and
TITLE OF INVENTION: Channel-1
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CURRENT FILING DATE: 2001-08-09
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TITLE OF INVENTION: Method and
TITLE OF INVENTION: Channel-1
                                                                           APPLICANT:
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APPLICANT: McSwiggen, Jim
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NUMBER OF SEQ ID NOS: 5450
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                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ENGTH: 17
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INVENTION: Method and Reagent for the Inhibition of Calcium Activated Chlori
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Thompson, Jim
McKenzie, Tim
                                                                    McSwiggen, Jim
Thompson, Jim
McKenzie, Tim
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Grupe, Andrew
                  Ayers, Dave
Grupe, Andrew
Szymkowski, Edmund
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100.0%; Pred. No.
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. 4.7e+02;
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; LENGTH: 17
; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-163-552-616
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US-10-163-552-616
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CURRENT APPLICATION NUMBER: US/10/251,117
CURRENT FILING DATE: 2003-02-24
PRIOR APPLICATION UNMEER: US 60/393,924
PRIOR FILING DATE: 2002-07-03
PRIOR APPLICATION NUMBER: US 10/163,552
PRIOR FILING DATE: 2002-06-06
PRIOR APPLICATION NUMBER: US 60/358,580
PRIOR APPLICATION NUMBER: US 60/358,580
PRIOR PRIOR DATE: 2002-02-20
PRIOR FILING DATE: 2002-02-20
PRIOR APPLICATION NUMBER: US 99/916,466
PRIOR APPLICATION NUMBER: US 60/296,249
PRIOR APPLICATION NUMBER: US 60/296,249
                                                                                                                                                                                                                                                                                                                                                    Sequence 147, Application US/10251117 Publication No. US20030170891A1 GENERAL INFORMATION:
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FILE REFERENCE: 249/021
CURRENT APPLICATION NUMBER: US/09/927,046
CURRENT FILING DATE: 2001-08-09
NUMBER OF SEQ ID NOS: 5450
SOFTWARE: PatentIn version 3.0
SEQ ID NO 305
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Publication No. US20030105051A1
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APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Nucleic acid treatment of diseases or conditions related to leve
TITLE OF INVENTION: HER2
FILE REFERENCE: MRHBH01-1653-A (400/014)
CURRENT APPLICATION NUMBER: US/10/163,552
CURRENT FILING DATE: 2002-06-06
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NUMBER OF SEQ ID NOS: 1997
SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                            TITLE OF INVENTION: RNA Interference Mediated Inhibition of Epidermal Growth Factor TITLE OF INVENTION: Gene Expression Using Short Interfering RNA FILE REFERENCE: 900/042 (MBHB02-468-A)
                                                                                                                                                                                                                                                                                                        APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: RNA
ORGANISM: Homo sapiens
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Similarity 84.6%;
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Similarity 100.0%;
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Pred. No.
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SEQ ID NOS:

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US-10-435-696-217; Sequence 217, Application US/10435696; Publication No. US20040018525A1
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CURRENT FILING DATE: 2003-02-24
PRIOR APPLICATION NUMBER: US 60/393,924
PRIOR FILING DATE: 2002-07-03
PRIOR APPLICATION NUMBER: US 10/163,552
PRIOR TILING DATE: 2002-06-06
PRIOR APPLICATION NUMBER: US 60/358,580
PRIOR APPLICATION NUMBER: US 60/358,580
PRIOR FILING DATE: 2002-02-20
PRIOR FILING DATE: 2002-02-20
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; SEQ ID NO 147
                                                                                                                        GENERAL INFORMATION:
APPLICANT: Wirtz, Ralph
APPLICANT: Munnes, Marc
APPLICANT: Kallabis, Ha
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn version 3.0 SEQ ID NO 396
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                   APPLICANT: Munnes, Marc
APPLICANT: Kallabis, Harald
APPLICANT: Kallabis, Harald
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR THE PREDICTION, DIAC
TITLE OF INVENTION: PREVENTION AND TREATMENT OF MALIGNANT NEOPLASIA
FILE REFERENCE: LeA 36 108
CURRENT APPLICATION NUMBER: US/10/435,696
CURRENT FILING DATE: 2003-05-09
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TITLE OF INVENTION: RNA Interference Mediated Inhibition of Epidermal Growth Factor
TITLE OF INVENTION: Gene Expression Using Short Interfering RNA
FILE REFERENCE: 900/042 (MBHB02-468-A)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US 09/916,466 PRIOR FILING DATE: 2001-07-25
PRIOR APPLICATION NUMBER: EP03003112.4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
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Pred. No.
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Pred. No. 5.9e+03;
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. 5.9e+03;
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CURRENT APPLICATION NUMBER: US/10/092,900A CURRENT FILING DATE: 2002-03-07
PRIOR APPLICATION NUMBER: USSN 60/274,322
PRIOR FILLING DATE: 2001-03-08
PRIOR APPLICATION NUMBER: USSN 60/283,675
PRIOR FILLING DATE: 2001-04-13
PRIOR APPLICATION NUMBER: USSN 60/338,092
PRIOR FILLING DATE: 2001-12-03
PRIOR FILLING DATE: 2001-12-03
PRIOR APPLICATION NUMBER: USSN 60/274,281
PRIOR APPLICATION NUMBER: USSN 60/274,281
PRIOR APPLICATION NUMBER: USSN 60/274,191
PRIOR FILLING DATE: 2001-03-08
PRIOR FILLING DATE: 2001-03-08
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US-10-092-900A-475
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; OTHER INFORMATION: LOC51242 for US-10-435-696-217
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Publication No. US20040043382A1
GENERAL INFORMATION;
APPLICANT: Padigaru, Muralidhara
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SEQ ID NO 217
LENGTH: 20
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Best Local (
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TITLE OF INVENTION: No. US20040043382A1el Proteins and Nucleic Acids Encoding Same
FILE REFERENCE: 21402-290C
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PRIOR APPLICATION NUMBER: EPO2010291.9
PRIOR FILING DATE: 2002-05-21
NUMBER OF SEQ ID NOS: 314
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ORGANISM: ARTIFICIAL SEQUENCE
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Zhong, Haihong
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Malyankar, Uriel M.
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Vernet, Corine A.M.
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Gusev, Vladimir Y.
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ilarity 100.0%;
Conservative (
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Pred. No.
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PRIOR FILING DATE: 2001-07-10
PRIOR PILING DATE: 2001-07-10
PRIOR FILING DATE: 2001-03-30
PRIOR FILING DATE: 2001-03-30
PRIOR APPLICATION NUMBER: USSN 60/294,899
PRIOR PILING DATE: 2001-05-31
PRIOR APPLICATION NUMBER: USSN 60/287,424
PRIOR FILING DATE: 2001-04-30
Remaining Prior Application data removed - Sef NUMBER OF SEQ ID NOS: 768
SEQ ID NO 475
LENGTH: 20
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US-10-092-900A-481
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APPLICANT: Padigaru, Muralidhara
APPLICANT: Spytek, Kimberly A.
APPLICANT: channy. Suresh G.
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APPLICANT: Burgess, Catherine E.
TITLE OF INVENTION: No. US20040043382Alel Proteins and Nucleic Acids Encoding
FILE REFERENCE: 21402-290C
CURRENT APPLICATION NUMBER: US/10/092,900A
CURRENT FILING DATE: 2002-03-07
PRIOR APPLICATION NUMBER: USSN 60/274,322
PRIOR APPLICATION NUMBER: USSN 60/274,322
PRIOR PILING DATE: 2001-03-08
PRIOR PILING DATE: 2001-03-08
PRIOR FILING DATE: 2001-03-08
PRIOR FILING DATE: 2001-04-13
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Taupier Jr., Raymond
Pena, Carol E.A.
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Tchernev, Velizar T.
Fernandes, Elma R.
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Gangolli, Esha A.
Vernet, Corine A.M.
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Alsobrook, John P.
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Gusev, Vladimir Y.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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5.9e+03;
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                                                                                                                                        US-10-333-108-26
                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/10/33,108
CURRENT FILING DATE: 2003-10-02
PRIOR APPLICATION NUMBER: PCT/EP01/07524
PRIOR FILING DATE: 2001-07-02
PRIOR APPLICATION NUMBER: EP 00115353.5
PRIOR FILING DATE: 2000-07-14
NUMBER OF SEQ ID NOS: 56
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                   Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Remaining Prior A
NUMBER OF SEQ ID
SEQ ID NO 481
                                                                                                                                                                                                          SEQ ID NO 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 26, Application US/10333108 Publication No. US20040076968A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Method For Detecting Pre-Disposition TITLE OF INVENTION: Hepatotoxicity FILE REFERENCE: 22140-1026
                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: F.
                                                                                                                                                     LENGTH: 20
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
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                                                                                   Local
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APPLICATION NUMBER: USSN 60/287,424
FILING DATE: 2001-04-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 2001-09-27
APPLICATION NUMBER: USSN 60/304,354
FILING DATE: 2001-07-10
APPLICATION NUMBER: USSN 60/279,995
FILING DATE: 2001-03-30
APPLICATION NUMBER: USSN 60/294,899
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: USSN 60/274,281
FILING DATE: 2001-03-08
APPLICATION NUMBER: USSN 60/274,191
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APPLICATION NUMBER: USSN 60/325,681
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                                  104 CCCTGTGTCCTAC 116
17
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Acuna, Gonzalo
Foernzler, Dorothee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hoffmann-La Roche AG
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                                                                     Conservative
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                                                             100.0%; **
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                                                                   Score 13; DB 16;
; Pred. No. 5.9e+03;
0; Mismatches 0;
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                                                                                                    Length 20;
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US-10-751-736-10681/c ; Sequence 10681, Application US/10751736 ; Publication No. US20040265230A1

RESULT 13

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RESULT 15
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; ORGANISM: homo sapiens
US-10-751-736-10681
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-751-736-10682/c
                                                          Sequence 42570, Application US/10098263B Publication No. US20030104410A1 GENERAL INFORMATION:
APPLICANT: Mittman, Michael TITLE OF INVENTION: Human Microarray FILE REFERENCE: 3118.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 100.0%;
Marches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: RNA
ORGANISM: RNAi
S-10-751-736-10682
                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn version 3.2
SEQ ID NO 10682
LENGTH: 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn version 3.2 SEQ ID NO 10681
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Brown, Eugene
APPLICANT: Liu, Wei
APPLICANT: Liu, Wei
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING COLON
TITLE OF INVENTION: CANCERS
FILE REFERENCE: AM100927 (031896-002000)
CURRENT APPLICATION NUMBER: US/10/751,736
CURRENT FILLING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US Provisional Application 60/438,000
PRIOR FILING DATE: 2003-01-06
NUMBER OF SEQ ID NOS: 54873
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Liu, Wei
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING COLON
TITLE OF INVENTION: CANCERS
FILE REFERENCE: AM100927 (031896-002000)
CURRENT APPLICATION NUMBER: US/10/751,736
CURRENT FILING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US Provisional Application 60/438,000
PRIOR FILING DATE: 2003-01-66
NUMBER OF SEQ ID NOS: 54873
                 CURRENT APPLICATION NUMBER: US/10/098,263B CURRENT FILING DATE: 2003-01-08
PRIOR APPLICATION NUMBER: 60/276,759
                                                                                                                                                                                                                                                                        14 TGCCAGGACCTGC 2
                                                                                                                                                                                                                                                                                                               13 TGCCAGGACCTGC 25
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L00.0%; Pred. No.
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100.0%; Pred. No. 5.9e+03;
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                                                                                                                                                                                                                                                                                                                                                                          DB 18; 1
                                                                                                                                                                                                                                                                                                                                                                                           Length 21;
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APPLICANT: Treacy, Maurice
APPLICANT: Treacy, Maurice
APPLICANT: Treacy, Maurice
APPLICANT: Bowman, Michael R.
APPLICANT: Spaulding, Vikki
APPLICANT: Spaulding, Vikki
APPLICANT: Carlin-Duckett, McKeough
APPLICANT: Kelleher, Kerry S.
APPLICANT: Kelleher, Kerry S.
APPLICANT: Conetice Institute, Inc.
TITLE OF INVENTION: SECRETED PROTEINS AND POLYNUCLEOTIDES ENCODING THEM
FILE REFERENCE: GI 6000-10A
CURRENT APPLICATION NUMBER: US/10/114,893
CURRENT FILING DATE: 2002-04-02
EARLIER APPLICATION NUMBER: 09/413,232
EARLIER FILING DATE: 1999-10-06
NUMBER OF SEQ ID NOS: 321
SOOTWARE: Patentin Ver. 2.0
SEQ ID NO 302
LENGTH - 24
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US-09-263-959-62/c
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; NAME/KBY: misc feature
; LOCATION: (2) —
; OTHER INFORMATION: biotinylated phosphoaramidite residue
US-10-114-893-302
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US-10-098-263B-42570
                             Sequence 62, Application US/09263959
Patent No. US20020150891A1
GENERAL INFORMATION:
APPLICANT: Hood, Leroy E.
APPLICANT: Rowen, Lee
APPLICANT: Koop, Ben F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 131066
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 42570
                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local :
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Publication No. US20020193567A1
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APPLICANT: McCoy, John M.
APPLICANT: LaVallie, Edward R.
APPLICANT: Collins-Racie, Lisa
APPLICANT: Evans, Cheryl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 2001-03-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
OTHER INFORMATION: oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 29
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APPLICANT: Rowen, Lee
APPLICANT: KOOP, Ben F.
TITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH UTI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                       Local Similarity
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100.0%; Pred. No. 5.7
ive 0; Mismatches
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100.0%; Pred. No.
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5.8e+03
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CORRESPONDENCE ADDRESS: ADDRESSEE: Seed and

STREET: 6300 C

6300 Columbia Center, 701 Fifth Avenue

Seed and Berry LLP

NUMBER OF SEQUENCES:

1279

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CURRENT FILING DATE: 2001-05-25;
PRIOR APPLICATION NUMBER: US 60/207,456;
PRIOR APPLICATION NUMBER: US 60/207,456;
PRIOR PRIOR DATE: 2000-05-26
PRIOR PPLICATION NUMBER: US 60/236,359;
PRIOR PILING DATE: 2000-09-27;
PRIOR PILING DATE: 2001-01-30;
PRIOR APPLICATION NUMBER: PCT/US01/00666;
PRIOR APPLICATION NUMBER: PCT/US01/00667;
PRIOR PILING DATE: 2001-01-30;
PRIOR PILING DATE: 2001-01-30;
PRIOR PILING DATE: 2001-01-30;
PRIOR APPLICATION NUMBER: PCT/US01/00664;
PRIOR PILING DATE: 2001-01-30;
PRIOR APPLICATION NUMBER: PCT/US01/00669;
PRIOR APPLICATION NUMBER: PCT/US01/00669;
PRIOR APPLICATION NUMBER: PCT/US01/00669;
PRIOR PILING DATE: 2001-01-30;
PRIOR FILING DATE: 2001-01-30;
PRIOR FILING DATE: 2001-01-30;
PRIOR FILING DATE: 2001-01-30;
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Best Local
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APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: ACOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (206) 682-60: INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/263,959
FILING DATE: 05-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION: TELEPHONE: (206) 622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: McMasters, David D.
REGISTRATION NUMBER: 33,963
REFERENCE/DOCKET NUMBER: 920010.426C2
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Similarity 100.0%; Pred. No. 2.2e+04;
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RESULT 19
US-09-866-108-2182/c
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-2181
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
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PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR PRIOR APPLICATION NUMBER: PCT/US01/00664
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Best Local Similarity
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APPLICANT: CHEN, Wensbeng
APPLICANT: SHANKON, Mark
APPLICANT: SHANKON, Mark
FITTLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AECMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
CURRENT FILING DATE: 2001-05-25
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OR APPLICATION NUMBER: PCT/US01/00665
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00668
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00663
OR FILING DATE: 2001-01-30
OR FILING DATE: 2001-01-30
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00662
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00661
OR FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00669
FILING DATE: 2001-01-30
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5. US20020048800A1
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US-09-866-108-2182
                                                          SOFTWARE: Aeo
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PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Aeomica Sequence Listing Engine SEQ ID NO 2182
LENGTH: 17
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PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
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                                                                                                    NUMBER OF SEQ ID NOS: 1575
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CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
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ORGANISM: Homo sapiens
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ITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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mes 12; Conserv
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APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00668
FILING DATE: 2001-01-22
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APPLICATION NUMBER: PCT/US01/00665
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APPLICATION NUMBER: PCT/US01/00662
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FILING DATE: 2001-01-30
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. 2.1e+04;
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; LENGTH: 17
; TYPE: DNA
; ORGANISM: Homo s
US-09-866-108-2184
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESS
FILE REFERENCE: ABOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2001-01-30
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                                                                                                                                              SOFTWARE: Aeomica
SEQ ID NO 2184
         Matches
                                           Query Match
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
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PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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PRIOR FILING DATE: 2001-01-30
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PRIOR APPLICATION NUMBER: PCT/US01/00664
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o. US20020048800A1
       . Similarity 12; Conserv
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INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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PENN, Sharron
HANZEL, David
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       Conservative
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Pred. No.
     Mismatches
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2.1e+04;
                                   Length 17,
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   Indels
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GENERAL INFORMATION
APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
ADDITICANT: HANZEL, David K
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US-09-866-108-2186/c
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US-09-866-108-2185/c
                                                                                                                                                                                                                                                                                                                                                            ; LENGTH: 17
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-2185
                                                                                    Sequence 2186, Application Patent No. US20020048800A1 GENERAL INFORMATION:
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Best Local
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PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
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CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
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FILING DATE: 2001-01-30
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12; Conserv
                                                                                                                                                                                                                  CCAGGACCTGCT
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PENN, Sharron (
HANZEL, David)
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RESULT 24
US-09-825-805-442
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-2186
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SOFTWARE: Aeomica Sequence
SEQ ID NO 2186
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APPLICANT: Beigelman, Leo
APPLICANT: Beaudry, Amber
APPLICANT: Karpeisky, Alex
APPLICANT: Adamic, Jasenka Matulic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
                                                                 APPLICANT: Sweedler, Dave
APPLICANT: Zinnen, Shawn
TITLE OF INVENTION: Nucleotide Triphosphate
FILE REFERENCE: MBHB00-831-F (400/009)
CURRENT APPLICATION NUMBER: US/09/825,805
CURRENT FILING DATE: 2001-09-27
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PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
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CURRENT FILING DATE: 2001-05-25
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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OR FILING DATE: 2000-09-27
OR APPLICATION NUMBER: PCT/US01/00666
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00667
OR APPLICATION NUMBER: PCT/US01/00667
OR APPLICATION NUMBER: PCT/US01/00664
OR APPLICATION NUMBER: PCT/US01/00664
OR FILING DATE: 2001-01-30
NT FILING DATE: 2001-09-27
APPLICATION NUMBER: 09/578,223
FILING DATE: 2000-05-23
APPLICATION NUMBER: 09/476,387
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APPLICATION NUMBER: PCT/US01/00670
FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00662
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00661
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00665
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                                                                                                                                                                                                                                                                                                                      INFORMATION:
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o. US20030004122A1
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US-09-927-046-906/c
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; ORGANISM: Homo sapiens
US-09-825-805-442
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US-09-927-046-1303/c
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CURRENT APPLICATION NUMBER: US/09/927,046
CURRENT FILING DATE: 2001-08-09
NUMBER OF SEQ ID NOS: 5450
SOFTWARE: PatentIn version 3.0
SEQ ID NO 906
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SEQ ID NO 442
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Best Local
                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                               Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Ayers, Dave—APPLICANT: Grupe, Andrew APPLICANT: Szymkowski, Edmund TITLE OF INVENTION: Method and Reagent for the Inhibition of Calcium Activated Chlor. TITLE OF INVENTION: Channel-1
                                                       APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: McSwiggen, Jim
APPLICANT: Thompson, Jim
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APPLICANT: McSwiggen, Jim
APPLICANT: Thompson, Jim
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PRIOR FILING DATE: 1998-11-04
                                        APPLICANT:
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         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                               TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 17
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APPLICATION NUMBER: 09/301,511
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APPLICATION NUMBER: 09/474,432
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McSwiggen, Jim
Thompson, Jim
McKenzie, Tim
Ayers, Dave
Grupe, Andrew
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Thompson, Jim
McKenzie, Tim
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Pred. No. 2.1e+04;
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US-09-740-332-3321/c
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US-09-927-046-1303
                                                                                                                                                            Sequence 3321, Application US/09740332
Publication No. US20030125270A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relat
TITLE OF INVENTION: Hepatitis C Virus Infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/740,332
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9704
SOPTWARE: PATENTIN VERSION 3.0
SEQ ID NO 1234
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Publication No. US20030125270A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment
TITLE OF INVENTION: Hepatitis C Virus Infection
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SEQ ID NO 1303
LENGTH: 17
                                                  NUMBER OF SEQ ID NOS: 9704
SOFTWARE: PatentIn version
SEQ ID NO 3321
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                                                                                                            FILE REFERENCE: RPI 400/003
CURRENT APPLICATION NUMBER: US/09/740,332
CURRENT FILING DATE: 2001-03-26
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OTHER INFORMATION: oligonucleotide substrate
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TYPE: RNA
ORGANISM: artificial sequence
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US-09-817-879-1234
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Publication No. US20030171311A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Hepatitis C virus Infection
FILE REFERENCE: MBHB00-801-F
CURRENT APPLICATION NUMBER: US/09/817,879
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9703
SOFTWARE: PatentIn version 3.0
SEQ ID NO 1234
LENGTH: 17
                                                                                                                                                                                  FILE REFERENCE: MBHB00-801-F
CURRENT APPLICATION NUMBER: US/09/817,879
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9703
SOFTWARE: PatentIn version 3.0
SEQ ID NO 3321
LENGTH: 17
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Publication No. US20030171311A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment
TITLE OF INVENTION: Hepatitis C Virus Infection
                                   Query Match
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ORGANISM: artificial sequence
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     Local Similarity 100 nes 12; Conservative
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; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-163-552-617
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US-10-163-552-617
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                                                                                                                                                                   PRIOR APPLICATION NUMBER: PCT/US02/09187
PRIOR FILING DATE: 2002-03-26
PRIOR APPLICATION NUMBER: US 60/296,876
PRIOR FILING DATE: 2001-06-08
PRIOR PPLICATION NUMBER: US 60/335,059
PRIOR PILING DATE: 2001-10-24
PRIOR APPLICATION NUMBER: US 60/337,055
PRIOR APPLICATION NUMBER: US 60/337,055
PRIOR APPLICATION NUMBER: US 60/358,580
PRIOR APPLICATION NUMBER: US 60/358,580
PRIOR PILING DATE: 2002-02-20
PRIOR FILING DATE: 2002-02-20
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SOFTWARE: PatentIn version 3.0
SEQ ID NO 617
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Best Local Similarity 83.3%;
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TITLE OF INVENTION: OLICONUCLEOTIDE MEDIATED
TITLE OF INVENTION: VIRUS REPLICATION
FILE REFERENCE: 400/042US (MEHB02-249-E)
CURRENT APPLICATION NUMBER: US/10/669,841
CURRENT FILING DATE: 2003-09-23
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APPLICANT: Lawrence, Blatt
APPLICANT: Dennis, Macejak
APPLICANT: James, McSwiggen
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APPLICATION NUMBER: US 09/740,332
FILING DATE: 2000-12-18
APPLICATION NUMBER: US 09/611,931
FILING DATE: 2000-07-07
APPLICATION NUMBER: US 09/504,321
                                                                                           APPLICATION NUMBER: US 60/363,124
FILING DATE: 2002-03-11
APPLICATION NUMBER: US 09/817,879
FILING DATE: 2001-03-26
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Kenneth, Draper
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Pred. No. 2.1e+04;
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                                                                                                               SOFTWARE: PatentIn version 3.0 SEQ ID NO 5914 LENGTH: 17
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SOFTWARE: PatentIn version 3.0
SEQ ID NO 3827
LENGTH: 17
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TITLE OF INVENTION: OLIGONUCLEOTIDE MEDIATED INHIBITION OF HEPATITIS B VIRUS AND HEPATITIE OF INVENTION: VIRUS REPLICATION
FILE REFERENCE: 400/042US (MBHB02-249-E)
CURRENT APPLICATION NUMBER: US/10/669,841
CURRENT FILING DATE: 2003-09-23
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PRIOR FILING DATE: 2002-03-26
PRIOR APPLICATION NUMBER: US 60/296,876
PRIOR FILING DATE: 2001-06-08
PRIOR APPLICATION NUMBER: US 60/335,059
PRIOR APPLICATION NUMBER: US 60/335,059
PRIOR APPLICATION NUMBER: US 60/335,059
PRIOR PILING DATE: 2001-10-24
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OTHER INFORMATION: oligonucleotide substrate
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ORGANISM: Artificial Sequence
      FEATURE:
OTHER INFORMATION: Description of Artificial Sequence:
                                                                     ORGANISM: Artificial Sequence
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OTHER INFORMATION: Description of Artificial Sequence:
                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 60/363,124
FILING DATE: 2002-03-11
APPLICATION NUMBER: US 09/817,879
FILING DATE: 2001-03-26
APPLICATION NUMBER: US 09/740,332
                                                                                                                                                                                                                                FILING DATE: 2000-02-15
                                                                                                                                                                                                                                                   APPLICATION NUMBER: US 09/504,321
                                                                                                                                                                                                                                                                       FILING DATE: 2000-12-18
APPLICATION NUMBER: US 09/611,931
FILING DATE: 2000-07-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US 60/358,580
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 60/337,055 FILING DATE: 2001-12-05
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Kenneth, Draper
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Dennis, Macejak
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Lawrence,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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Pred. No. 2.1e+04;
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                                                                                                                                                                                                    See File Wrapper or PALM
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                           Nucleic Acid
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TITLE OF INVENTION: HUMAN MYOSIN-LIKE POLYPEPTIDE EXPRESSED PREDOMINANTLY IN HEART AN
FILE REFERENCE: PB0105
CURRENT APPLICATION NUMBER: US/10/723,361
CURRENT PILING DATE: 2003-11-26
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-010-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
RESULT 35

US-10-723-361-2182/c

; Sequence 2182, Application US/10723361

; Publication No. US20040137589A1

; GENERAL INFORMATION:
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Best Local Similarity
Tatches 12; Conserv:
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Best Local S
Matches 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Remaining Prior Application data removed NUMBER OF SEQ ID NOS: 15755
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APPLICANT: JI, Yongga
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PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
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                                                                                                                                                                                                                                                                     Local Similarity
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N, Sharron
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100.0%; Pred. No. 2.1e+04;
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Pred. No.
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2.1e+04;
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GU, Yizhong

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US-10-723-361-2183/c
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US-10-723-361-2182
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Best Local Similarity
APPLICANT: RANK, DAVIG R.

APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, MAIK
TITLE OF INVENTION: HUMAN MYOSIN-LIKE POLYPEPTIDE EXPRESSED PREDOMINANTLY IN HEART AN
FILE REFERENCE: PB0105
CURRENT APPLICATION NUMBER: US/10/723,361
CURRENT FILING DATE: 2003-11-26
PRIOR APPLICATION NUMBER: US 09/866,108
PRIOR FILING DATE: 2001-05-25
PRIOR FILING DATE: 2001-05-26
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-010-04
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-010-04
PRIOR APPLICATION NUMBER: GB 24263.5
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APPLICANT: JI, Yonggar
APPLICANT: PENN, Shari
APPLICANT: HANZEL, Dav
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PRIOR FILING DATE: 2001-01-30
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PRIOR FILING DATE: 2000-05-26
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PRIOR APPLICATION NUMBER: US 09/866,108
PRIOR FILING DATE: 2001-05-25
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CURRENT APPLICATION NUMBER: US/10/723,361
CURRENT FILING DATE: 2003-11-26
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TITLE OF INVENTION: HUMAN MYOSIN-LIKE POLYPEPTIDE EXPRESSED PREDOMINANTLY IN HEART
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FILING DATE: 2001-01-30
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PENN, Sharron G.
HANZEL, David K.
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PENN, Sharron G.
HANZEL, David K.
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Pred. No.
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RESULT 37
US-10-723-361-2184/c
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US-10-723-361-2183
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                                                                                                                                              PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
PRIOR PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR APPLICATION NUMBER: PCT/US01/00668
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PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
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PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
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CURRENT FILING DATE: 2003-11-26
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                                                                                     SOFTWARE: Aeomica Sequence
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TITLE OF INVENTION: HUMAN MYOSIN-LIKE POLYPEPTIDE EXPRESSED PREDOMINANTLY IN HEART A
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PRIOR FILING DATE: 2001-01-30
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ORGANISM: Homo sapiens
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APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
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                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: PCT/US01/00667 FILING DATE: 2001-01-30
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                                                                                                                              Prior Application data removed -
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RANK, David R.
CHEN, Wensheng
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                                                                                     Listing Engine
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Pred. No.
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RESULT 39
US-10-723-361-2186/c
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US-10-723-361-2185/c
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                                   APPLICANT: GU, Yizhong
                                                                      Sequence 2186, Application US/10723361
Publication No. US20040137589A1
                                                                                                                                                                                                                                                          Matches
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Best Local
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PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-01
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
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PRIOR TLING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
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CURRENT APPLICATION NUMBER: US/10/723,361
CURRENT FILLING DATE: 2003-11-26
PRIOR APPLICATION NUMBER: US 09/866,108
PRIOR FILING DATE: 2001-05-25
PRIOR FILING DATE: 2001-05-25
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TITLE OF I
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APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron
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PRIOR FILING DATE: 2000-05-26
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12; Conserv
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INVENTION: HUMAN MYOSIN-LIKE POLYPEPTIDE EXPRESSED PREDOMINANTLY IN HEART
                                                                                                                                                                                                                                                                                                                                                                                                                 Aeomica Sequence Listing Engine
                                                                                                                                                                                  CCAGGACCTGCT 2
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JI, Yonggang
PENN, Sharron G.
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HANZEL, David K.
                                                                                                                                                                                                                                                      Conservative
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100.0%; Pred. No.
                                                                                                                                                                                                                                                Score 12; DB 17; ; Pred. No. 2.1e+04; 0; Mismatches 0;
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Query Match
Best Local Similarity
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US-10-723-361-2186
                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 3339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 3339, Application US/09969373 Patent No. US20020133852A1 GENERAL INFORMATION:
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SEQ ID NO 2186
LENGTH: 17
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PRIOR FILING DATE: 2001-05-15
NUMBER OF SEQ ID NOS: 4593
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/969,373
CURRENT FILING DATE: 2001-10-02
PRIOR APPLICATION NUMBER: US 09/754,853
PRIOR FILING DATE: 2001-01-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Effertz, Roger J.
APPLICANT: Hauge, Brian M.
TITLE OF INVENTION: Soybean SSRs and Methods of Genotyping
FILE REFERENCE: 38-10(52679)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 09/760,427 PRIOR FILING DATE: 2001-01-13
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PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2000-09-27
                                                                                                                                                                                                                             TYPE: DNA ORGANISM: Glycine
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
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PRIOR APPLICATION NUMBER: PCT/US01/00668
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PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
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CURRENT FILING DATE: 2003-11-26
CURRENT APPLICATION NUMBER: US 09/866,108
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ORGANISM:
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00667
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
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INVENTION: HUMAN MYOSIN-LIKE POLYPEPTIDE EXPRESSED PREDOMINANTLY IN HEART AND ADDRESSED PREDOMINANTLY AND ADDRESSED PREDOMINANTLY ADDRESSED PREDOMINANT AND ADDRESSED PREDOMINANT ADDRESSED PREDOMINANT AND ADDRESSED PREDOMINANT AND ADDRESSED PREDOMINANT ADDRESSED PREDOMINANT AND ADDRESSED PREDOMINANT AND ADDRESSED PREDOMINANT AND ADDRESSED PREDOMINANT ADDRESSED PREDOMINANT AND ADDRESSED PREDOMINAN
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RANK, David R.
CHEN, Wensheng
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                       9.1%; 5c.
100.0%; Pr
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                           ore 12; DB 9; Lored. No. 2.1e+04; Mismatches 0;
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2.1e+04;
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Conservative

Indels

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Gaps

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; SOFTWARE: PatentIn veri
; SEQ ID NO 22
; LENGTH: 20
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-767-421-22
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US-09-767-421-22
; Sequence 22, Application US/09767421
; Publication No. US20030175954A1
; GENERAL INFORMATION:
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SEQ ID NO 110
LENGTH: 19
TYPE: DNA
                                                                                                     Query Match
Best Local Similarity
Matches 12; Conserv
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Patent No. US20020142359A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Edge, Michael Derek
APPLICANT: Emery, Stephen Charles
TITLE OF INVENTION: Monoclonal Antibody to CEA, Conjugates Comprising Said Antibody,
TITLE OF INVENTION: Monoclonal Antibody to CEA, Conjugates Comprising Said Antibody,
FILE REFERENCE: 1991-209
FILE REFERENCE: 1991-209
CURRENT APPLICATION NUMBER: US/09/910,059
CURRENT FILING DATE: 1998-10-29
PRIOR APPLICATION NUMBER: US 09/171,945
PRIOR APPLICATION NUMBER: PCT/GB97/01165
PRIOR PILING DATE: 1997-04-29
PRIOR APPLICATION NUMBER: GB 9703103.3
PRIOR FILING DATE: 1997-02-14
PRIOR APPLICATION NUMBER: GB 9703103.3
PRIOR FILING DATE: 1997-02-14
PRIOR APPLICATION NUMBER: GB 9703103.7
PRIOR FILING DATE: 1996-05-04
NUMBER: OF SEQ ID NOS: 131
NUMBER: OF SEQ ID NOS: 131
                                                                                                                                                                                                                                                                                                         FILE REFERENCE: Unassigned CURRENT APPLICATION NUMBER: US/09/767,421 CURRENT FILING DATE: 2001-01-22 NUMBER OF SEQ ID NOS: 74
                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Shamblott, Michael
APPLICANT: Gearhart, John
TITLE OF INVENTION: Human Embryoid Body-Derived Cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Copley, Clive G
APPLICANT: Edge, Michael Derek
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mes 12; Conserv
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                         CCAGGACCTGCT 13
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                                                                                                                        9.1%; Score 12;
100.0%; Pred. No.
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100.0%; Pred. No.
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2.1e+04;
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US-10-175-239-15/c
; Sequence 15, Application US/10175239
; Publication No. US20030232774A1
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; OTHER INFORMATION: antisense sequence
US-10-238-442-99
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US-10-238-442-99:
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US-10-161-996-81/c
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CURRENT FILING DATE: 2002-09-09
PRIOR APPLICATION NUMBER: 09/640,101
PRIOR FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: 09/286,904
PRIOR APPLICATION NUMBER: 09/286,904
PRIOR FILING DATE: 1999-04-06
NUMBER OF SEQ ID NOS: 107
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                           US-10-161-996-81
                                                                           Query Match
Best Local S
Matches 12
                                                                                                                                                                         NUMBER OF SEQ ID NOS: 273

SEQ ID NO 81
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE: OTHER INFORMATION: Antisense Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 81, Application US/10161996 Publication No. US20030224515A1
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Publication No. US20030176383A1
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Best Local Similarity
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APPLICANT: Nero, Pamela S.
APPLICANT: MCKAy, Pamela S.
TITLE OF INVENTION: Antisense Modulation of p38 Mitogen
TITLE OF INVENTION: Activated Protein Kinase Expression
FILE REFERENCE: ISPH-0488
                                                                                                                                                                                                                                                                                                                    APPLICANT: Brenda F. Baker
APPLICANT: Kenneth W. Dobie
TITLE OF INVENTION: ANTISENSE MODULATION OF STEROL REGULATORY ELEMENT-BINDING PROTEI
TITLE OF INVENTION: EXPRESSION
FILE REFERENCE: RTS-0395
CURRENT APPLICATION NUMBER: US/10/161,996
CURRENT FILING DATE: 2002-06-04
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TYPE: DNA
ORGANISM: Artificial Sequence
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                                     24 GCTGCACGACTG 35
20 GCTGCACGACTG
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100.0%; Pred. No.
tive 0; Mismatc
                                                                           9.1%; Score 12; DB 15;
100.0%; Pred. No. 2.1e+04
tive 0; Mismatches 0
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hes 0;
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2.1e+04;
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RESULT 45

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; TYPE: DNA
; ORGANISM: Chlamydia
US-10-289-762-5153
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US-10-289-762-5153
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US-10-175-239-52
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                                                                                                                                                                                    Sequence 5153, Application US/10289762
Publication No. US20040006218A1
GENERAL INFORMATION:
APPLICANT: Griffals, R.
TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence and polypeptides,
TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnos
TITLE OF INVENTION: and treatment of infection
FILE REFERENCE: 9710-003-999
FILE REFERENCE: 9710-003-999
                                                                                            SEQ ID NO 5153
LENGTH: 20
   Query Match
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Best Local
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SEQ ID NO 52
LENGTH: 20
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Publication No. US20030232774A1
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LENGTH: 20
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Best Local
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TITLE OF INVENTION: ANTISENSE MODULATION OF PROFILIN 1 EXPRESSION
FILE REFERENCE: HTG-0017
CURRENT APPLICATION NUMBER: US/10/175,239
CURRENT FILING DATE: 2002-06-17
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APPLICANT: Kenneth W. Dobie
TITLE OF INVENTION: ANTISENSE MODULATION OF PROFILIN 1 EXPRESSION
FILE REFERENCE: HTS-0017
CURRENT APPLICATION NUMBER: US/10/175,239
CURRENT FILING DATE: 2002-06-17
NUMBER OF SEQ ID NOS: 79
                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/289,762
CURRENT FILING DATE: 2003-03-27
NUMBER OF SEQ ID NOS: 6849
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: H. sapiens
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ORGANISM: Artificial Sequence
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Score 12;
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RESULT 50
US-10-304-105-36/c
IS-quence 36, Application US/10304105
; Publication No. US20040101854A1
; GENERAL INFORMATION:
; APPLICANT: C. Frank Bennett
; APPLICANT: Nicholas M. Dean
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US-10-289-762-5261/c
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LENGTH: 20
TYPE: DNA
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CURRENT APPLICATION NUMBER: US/10/304,105
CURRENT FILING DATE: 2002-11-21
NUMBER OF SEQ ID NOS: 68
SEQ ID NO 30
LENGTH: 20
TYPE: DNA
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Publication No. US20040101854A1
GENERAL INFORMATION:
APPLICANT: C. Frank Bennett
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Publication No. US20040006218A1
GENERAL INFORMATION:
APPLICANT: Griffals, R.
APPLICANT: Griffals, R.
TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence and polypeptides, fragment:
TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnosis, previous of inventions and treatment of infection
                                                                                                                                                                                                                                             Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Nicholas M. Dean
APPLICANT: Kenneth W. Dobie
TITLE OF INVENTION: MODULATION OF BCL2-ASSOCIATED ATHANOGENE EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/289,762
CURRENT FILING DATE: 2003-03-27
NUMBER OF SEQ ID NOS: 6849
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                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Antisense Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Chlamydia pneumoniae
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100.0%; Pred. No. 2.1e+04
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100.0%; Pred. No.
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); Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                           RESULT 52
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Best Local S
Matches 12
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LENGTH: 20
                                                                                                                                                                                                                                                            Sequence 99, Application US/10641455A
Publication No. US20040171566A1
GENERAL INFORMATION:
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SEQ ID NO 36
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
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Best Local
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               APPLICANT: Gaarde, William A.
APPLICANT: Nero, Pamela S.
APPLICANT: McKay, Robert
APPLICANT: McKay, Robert
APPLICANT: Popoff, Ian
ITTLE OF INVENTION: Antisense Oligonucleotide Modulation of p38 Mitogen
TITLE OF INVENTION: Activated Protein Kinase Expression
FILE REFERENCE: ISPH-0762
CURRENT APPLICATION NUMBER: US/10/641,455A
CURRENT FILING DATE: 2003-08-15
PRIOR APPLICATION NUMBER: US 10/238,442
PRIOR FILING DATE: 2002-09-09
                                                                                                                                                                                                                      APPLICANT: Monia, Brett P. APPLICANT: Gaarde, Willia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: MODULATION OF THYROID HORMONE RECEPTOR INTERACTOR 3 EXPRESSION FILE REFERENCE: PTS-0076
CURRENT APPLICATION NUMBER: US/10/346,268
CURRENT FILING DATE: 2003-01-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: C. Frank Bennett
APPLICANT: Nicholas M. Dean
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/10/304,105
CURRENT FILING DATE: 2002-11-21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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les 12; Conserv
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APPLICATION NUMBER: US 09/640,101
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100.0%; Pred. No. 2.1e+04;
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100.0%; Pred. No.
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APPLICANT: Suean M. Freier
FITTLE OF INVENTION: ANTISENSE MODULATION OF SAP-1 EXPRESSION
FILE REFERENCE: RTS-0267
CURRENT APPLICATION NUMBER: US/10/484,669
CURRENT FILING DATE: 2004-01-23
PRIOR APPLICATION NUMBER: US/09/920,759
PRIOR APPLICATION NUMBER: US/09/920,759
PRIOR FILING DATE: 2001-08-01
NUMBER OF SEQ ID NOS: 91
SEQ ID NO 53
LENGTH: 20
TYPE: DNA
TYPE: DNA
TYPE: DNA
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US-10-484-669-53/c
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                                   SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 59 LENGTH: 21
                                                                                                                                                                                                                                                                                                    Sequence 59, Application US/10010066 Publication No. US20020173001A1
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Best Local Similarity
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                                                                                       APPLICANT: University of Iowa Research Foundation
APPLICANT: Schwartz, David A.
APPLICANT: Schwartz, David A.
APPLICANT: Schutte, Brian C.
TITLE OF INVENTION: Variant TLR4 nucleic acid and uses thereof
FILE REFERENCE: 875.010W01
CURRENT APPLICATION NUMBER: US/10/010,066
CURRENT FILING DATE: 2001-12-07
PRIOR APPLICATION NUMBER: US 09/329,515
PRIOR PILING DATE: 1999-06-10
NUMBER OF SEC ID NOS: 72
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                 LENGTH: 21
TYPE: DNA
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TYPE: DNA
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SOFTWARE: PatentIn Ver. 2.0
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OTHER INFORMATION: antisense sequence
ORGANISM: Homo sapiens
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100.0%; Pred. No.
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100.0%; Pred. No.
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. 2.1e+04;
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2.1e+04;
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APPLICANT: Gokarn, Ravi R
APPLICANT: Jessen, Holly
APPLICANT: Jessen, Holly
APPLICANT: Zidwick, Mary Jo
TITLE OF INVENTION: Isopremoid Production
FILE REFERENCE: 12904/002US1
CURRENT APPLICATION NUMBER: US/10/381,779
CURRENT FILING DATE: 2003-03-28
PRIOR APPLICATION NUMBER: PCT/US01/30328
PRIOR APPLICATION NUMBER: PCT/US01/30328
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-29
NUMBER OF SEQ ID NOS: 190
NUMBER OF SEQ ID NOS: 190
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US-10-381-779-183/c
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US-10-758-672A-29
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                                                                                   ; OTHER INFORMATION: Synthetic primer US-10-758-672A-29
                                                                                                                                                                                PRIOR APPLICATION NUMBER: US 60/187,911
PRIOR FILING DATE: 2000-03-08
NUMBER OF SEQ ID NOS: 29
SOFTWARE: PatentIn version 3.2
SEQ ID NO 29
LENGTH: 21
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SEQ ID NO 183
LENGTH: 21
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Best Local
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      Query Match 9.1%; Score 12; DB Best Local Similarity 100.0%; Pred. No. 2.3 Matches 12; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                 APPLICANT: HAN, et al.
TITLE OF INVENTION: HUMAN E3 ALPHA UBIQUITIN LIGASE FAMILY
FILE REFERENCE: 101017/35966B
CURRENT APPLICATION NUMBER: US/10/758,672A
CURRENT FILING DATE: 2004-01-15
PRIOR APPLICATION NUMBER: US 09/724,126
PRIOR FILING DATE: 2000-11-28
PRIOR FILING DATE: 2000-11-28
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                                                                                                                       TYPE: DNA
ORGANISM: Artificial sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
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; Pred. No.
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      DB 17; Length 21; 2.1e+04; hes 0; Indels
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          RESULT 59
US-10-697-036-83
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                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Serup, Palle
APPLICANT: Heimberg, Harry
APPLICANT: Gradwohl Gerard
TITLE OF INVENTION: Methods For Generating Insulin-Secreting
TITLE OF INVENTION: Cells Suitable for Transplantation
FILE REFERENCE: 6246.200-US
CURRENT APPLICATION NUMBER: US/10/090,011
CURRENT FILING DATE: 2002-02-26
PRIOR APPLICATION NUMBER: US 60/271,474
PRIOR FILING DATE: 2001-02-26
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                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 70
SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 5
LENGTH: 22
TYPE: DNA
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SEQ ID NO 10507
LENGTH: 21
                                                                       Matches
                                                                                         Query Match
Best Local Similarity
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CURRENT APPLICATION NUMBER: US/10/751,736
CURRENT FILING DATE: 2003-01-06
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING COLON
TITLE OF INVENTION: CANCERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
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ORGANISM: homo sapiens
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                                  6 CATCATCTGCCA 17
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CATCATCTGCCA 17
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                                                                       Conservative
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100.0%; Pred. No. 2.1e+04;
tive 0; Mismatches 0;
                                                                                          100.0%;
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                                                                                          Score 12; DB 14;
; Pred. No. 2.1e+04;
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                                                                                                          Length 22
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RESULT 61
US-09-779-880A-38/c
; Sequence 38, Application US/09779880A
; Patent No. US20020061834A1
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US-09-779-879A-38/c
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PRIOR FILING DATE: 2002-10-31
NUMBER OF SEQ ID NOS: 90
SOFTWARE: Patentin version 3.2
SEQ ID NO 83
LENGTH: 22
TYPE: DNA
TYPE: DNA
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Best Local
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APPLICANT: Ruben, Steven, M.
APPLICANT: Ruben, Steven, M.
TITLE OF INVENTION: Human G-protein Chemokine Receptor (CCR5) HDGNR10
FILE REFERENCE: 1488.115000A
CURRENT FILING DATE: 2001-02-09
FRIOR APPLICATION NUMBER: US 60/181,258
PRIOR PILING DATE: 2000-02-09
PRIOR PILING DATE: 2000-03-09
PRIOR PILING DATE: 2000-09-22
RUMBER OF SEQ ID NOS: 58
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TITLE OF INVENTION: TRANSFORMED CELL WITH ENHANCED SENSITIVITY TO ANTIFUNGAL COMPOUND
TITLE OF INVENTION: THEREOF
FILE REFERENCE: 078242
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Rosen,
APPLICANT: Roschi
APPLICANT: Li, Y
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CURRENT FILING DATE: 2003-10-31
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TYPE: DNA
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OTHER INFORMATION: Description OTHER INFORMATION: oligonucles
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OTHER INFORMATION: 5' Oligonu
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                                                                                                                                                                                                                                                                                                           5' Oligonucleotide primer
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100.0%; Pred. No. 2.1e+04;
tive 0; Mismatches 0;
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100.0%; Pred. No. 2.1e+04;
tive 0; Mismatches 0; Indels
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RESULT 63
US-09-910-120-26/c
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CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: US 60/181,258
PRIOR FILING DATE: 2000-02-09
PRIOR APPLICATION NUMBER: US 60/187,999
PRIOR APPLICATION NUMBER: US 60/234,336
PRIOR FILING DATE: 2000-09-22
PRIOR FILING DATE: 2000-09-22
NUMBER OF SEQ ID NOS: 58
COMMENCE: DESCRIPTION OF SETEMAN OF SEQ ID NOS: 58
                                                                                                                                                                                                                                                                                                                                              SEQ ID NO 26
LENGTH: 23
TYPE: DNA
Sequence 26, Application US/09910120 Patent No. US20020137053A1
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LENGTH: 23
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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Best Local
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TITLE OF INVENTION: NOVEL FAB FRAGMENT LIBRARIES AND METHOD FOR THEIR USE
FILE REFERENCE: DX/003 CON
CURRENT APPLICATION NUMBER: US/09/988,899
CURRENT FILING DATE: 2001-11-19
                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 99201558.6 PRIOR FILING DATE: 1999-05-18 NUMBER OF SEQ ID NOS: 71
                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn Ver. 2.1
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TITLE OF INVENTION: Human G-protein Chemokine Receptor
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                                                                                                                                                                                                                                                                                FEATURE: OTHER INFORMATION: Description of Artificial Sequence: Primer
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100.0%; Pr
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100.0%; Pred. No.
tive 0; Mismatc
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2.1e+04;
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hes 0;
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APPLICANT: DANA AULT-RICHE

INFORMATION

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RESULT 65
US-09-833-041-51/c
; Sequence 51, Application US/09833041
; Publication No. US20030125247A1
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US-09-939-769-45/c
; Sequence 45, Application US/09939769
; Publication No. US20030017149A1
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; OTHER INFORMATION: Primer: HuVkappa5aBACK
US-09-910-120-26
                                                                                                                                                                                                                                                                                                                                                                                                                      ; OTHER INFORMATION: Description of Artificial Sequence: Primer US-09-939-769-45
                                             GENERAL INFORMATION:
APPLICANT: Rosen, Craig A.
APPLICANT: Haseltine, William A.
TITLE OF INVENTION: Albumin Fusion Proteins
FILE REFERENCE: PF545
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 1996-10-10
NUMBER OF SEQ ID NOS: 96
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 45
LENGTH: 23
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PRIOR FILING DATE: 2000-07-19
NUMBER OF SEQ ID NOS: 73
SOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 26
LENGTH: 23
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Best Local
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CURRENT APPLICATION NUMBER: US/09/910,120
CURRENT FILING DATE: 2001-07-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: HOEFFLER, JAMES P.
APPLICANT: RUSSELL, MARIJANE
TITLE OF INVENTION: SINGLE CHAIN ANTIBODY FUSION REAGENTS THAT REGULATE
TITLE OF INVENTION: TRANSCRIPTION IN VIVO
FILE REFERENCE: 039322/0226
CURRENT FILLING DATE: 2001-08-28
CURRENT FILLING DATE: 2001-08-28
ENTOR APPLICATION NUMBER: 08/728,890
PRIOR APPLICATION NUMBER: 08/728,890
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TITLE OF INVENTION: COLLECTIONS OF BINDING PROTEINS AND TAGS
TITLE OF INVENTION: AND USES THEREOF FOR NESTED SORTING AND HIGH THROUGHPUT
TITLE OF INVENTION: SCREENING
                 CURRENT APPLICATION NUMBER: US/09/833,041
CURRENT FILING DATE: 2001-04-12
PRIOR APPLICATION NUMBER: 60/229,358
                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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ORGANISM: Artificial Sequence
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Pred. No.
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                                                                                             GENERAL INFORMATION:
                                                                                                                 Sequence 51, Application US/09832929 Publication No. US20040171123A1
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; OTHER INFORMATION: Degenerate Vkappa forward primer useful ; OTHER INFORMATION: amplifying human VL domains US-09-833-041-51
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US-09-832-929-51/c
                   RESULT 67
                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/833,245
CURRENT FILING DATE: 2001-04-12
PRIOR APPLICATION NUMBER: 60/229, 358
PRIOR FILING DATE: 2000-04-12
PRIOR APPLICATION NUMBER: 60/256, 931
PRIOR FILING DATE: 2000-12-21
PRIOR APPLICATION NUMBER: 60/199, 384
PRIOR PILING DATE: 2000-04-25
NUMBER OF SEQ ID NOS: 2267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 12; Conser
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LENGTH: 23
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                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 51
LENGTH: 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 51, Application US/09833245 Publication No. US20040010134A1
                                                                                                                                        Matches
                                                                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 2000-04-25
NUMBER OF SEQ ID NOS: 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR FILING DATE: 2000-04-12
PRIOR APPLICATION NUMBER: 60/256,931
PRIOR FILING DATE: 2000-12-21
PRIOR APPLICATION NUMBER: 60/199,384
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PEATURE: primer bind
                                                                                                                                                                                                                      OTHER INFORMATION: Degenerate Vkappa OTHER INFORMATION: amplifying human V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                            FEATURE:
NAME/KEY: primer_bind
                                                                                                                                                                                                                                                                                            TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
                                                                                                                                                        Local
                                                                                                      31 GACTGCGTGAGT 42
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                                                                     20 GACTGCGTGAGT 9
                                                                                                                                        l Similarity
12; Conserv
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100.0%; Pr
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Pred. No.
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. 2.1e+04;
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2.1e+04;
                                                                                                                                                                        Length 23
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APPLICANT: Rosen, Craig A.
APPLICANT: Haseltine, William A.
TITLE OF INVENTION: Albumin Fusion Proteins
FILE REFERENCE: PF547

CURRENT APPLICATION NUMBER: US/09/832,929

CURRENT FILING DATE:

2001-04-12

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PRIOR APPLICATION NUMBER: 60/229,358
PRIOR FILING DATE: 2000-04-12
PRIOR APPLICATION NUMBER: 60/256,931
PRIOR FILING DATE: 2000-12-21
PRIOR APPLICATION NUMBER: 60/199,384
PRIOR FILING DATE: 2000-04-25
NUMBER OF SEQ ID NOS: 72
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 51
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US-10-039-785-21/c
                                                                                                                           ; OTHER INFORMATION: PCR primer useful US-10-039-785-21
                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 66
SOFTWARE: PatentIn Ver. 2
SEQ ID NO 21
LENGTH: 23
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Publication No. US20020067646A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Salcedo et al.
                                             Matches
                                                               Query Match
Best Local :
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PRIOR APPLICATION NUMBER: 60/294,981
PRIOR FILING DATE: 2001-06-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/369,860
PRIOR FILING DATE: 2002-04-05
PRIOR APPLICATION NUMBER: 60/341,237
PRIOR FILING DATE: 2001-12-20
PRIOR APPLICATION NUMBER: 60/331,310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/10/039,785
CURRENT FILING DATE: 2002-05-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Antibodies that Immunospecifically Bind to TRAIL TITLE OF INVENTION: Receptors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 2001-11-07
PRIOR APPLICATION NUMBER: 60/327,364
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 2001-11-14
PRIOR APPLICATION NUMBER: 60/331,044
                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE: 2001-05-25
                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/293,473
                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/309,176
                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR FILING DATE: 2001-09-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/323,807
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TYPE: DNA
ORGANIEM: Artificial Sequence
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NAME/KEY: primer_bind
OTHER INFORMATION: Degenerate Vkappa forward primer useful
OTHER INFORMATION: amplifying human VL domains
                                                                                                                                                                                    TYPE: DNA
ORGANISM: Artificial sequence
                                                                                                                                                                     FEATURE:
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Local Similarity 100.0%;
nes 12; Conservative (
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                                         l Similarity
12; Conser
                                       9.1%; Score 12; DB 13; Length 23; Larity 100.0%; Pred. No. 2.1e+04; Conservative 0; Mismatches 0; Indels
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Pred. No.
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. 2.1e+04;
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SEQ ID NO 48
LENGTH: 23
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Degenerate Vkappa forward primer useful
OTHER INFORMATION: amplifying human VL domains
US-10-153-064-48
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US-10-153-064-48/c
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                                                                                                                                                                                                                  SEQ ID NO 100
LENGTH: 23
TYPE: DNA
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CURRENT FILING DATE: 2002-02-15
PRIOR APPLICATION NUMBER: 60/272,107
PRIOR FILING DATE: 2001-02-28
PRIOR APPLICATION NUMBER: 60/209,811
PRIOR FILING DATE: 2000-06-06
NUMBER OF SEQ ID NOS: 138
SOFTWARE: Patentin Ver. 2.1
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Publication No. US20020142814A1
GENERAL INFORMATION:
APPLICANT: Bell et al.
TITLE OF INVENTION: Chemokine Beta-1 Fusion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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                                                                           Matches
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                                                                                             Best Local Similarity
                                                                                                          Query Match
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CURRENT APPLICATION NUMBER: US/10/153,064
CURRENT FILING DATE: 2002-05-24
PRIOR APPLICATION NUMBER: 60/293,212
PRIOR FILING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 137
                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: ZHOU, XIA-DI
APPLICANT: PEACH, ROBERT
TITLE OF INVENTION: B7-RELATED NUCLEIC
TITLE OF INVENTION: IMMUNOMODULATION
FILE REFERENCE: 3053-4071US3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                               OTHER INFORMATION: Description of Artificial Sequence: Primer
                                                                                                                                                                                ORGANISM: Artificial Sequence FEATURE:
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                                      31 GACTGCGTGAGT 42
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Il Similarity 100.0%; F
12; Conservative 0;
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YANG, GUCHEN
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                                                                           Conservative
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                                                                                         9.1%; Score 12;
100.0%; Pred. No.
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Pred. No.
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; ORGANISM: Homo sapiens
US-10-056-884-46
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LENGTH: 23
Sequence 43, Application US/10080980
Publication No. US20030036115A1
GENERAL INFORMATION:
APPLICANT: Bristol-Myers Squibb Company
TITLE OF INVENTION: POLYNUCLECTIDE ENCODING A NOVEL HUMAN POTASSIUM CHANNEL BETA-SUBU
TITLE OF INVENTION: K-betam6, EXPRESSED HIGHLY IN THE SMALL INTESTINE
FILE REFERENCE: D0121 NP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 46, Application US/10056884 Publication No. US20030032786A1 GENERAL INFORMATION:
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LENGTH: 23
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Best Local
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Bristol-Myers Squibb Company
TITLE OF INVENTION: FOLYNUCLECTIDE ENCODING A NOVEL HUMAN FOTASSIUM CHANNEL BETA-SUBU
TITLE OF INVENTION: K-betaM2
FILE REFERENCE: DOO76 NP
CURRENT APPLICATION NUMBER: US/10/056,884
CURRENT FILING DATE: 2002-01-24
PRIOR APPLICATION NUMBER: US 60/263,872
PRIOR FILING DATE: 2001-01-24
PRIOR FILING DATE: 2001-01-24
PRIOR FILING DATE: 2001-02-14
NUMBER OF SEQ ID NOS: 73
NUMBER OF SEQ ID NOS: 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: D0079 NP
CURRENT APPLICATION NUMBER: US/10/075,846
CURRENT FILING DATE: 2002-02-13
PRIOR APPLICATION NUMBER: US 60/269,535
PRIOR FILING DATE: 2001-02-16
NUMBER OF SEQ ID NOS: 81
SOFTWARE: Patentin version 3.0
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TITLE OF INVENTION: POLYNUCLEOTIDES ENCODING A NOVEL GLYCINE RECEPTOR ALPHA SUBUNIT I
TITLE OF INVENTION: IN THE GASTROINTESTINAL TRACT, HGRA4, and SPLICE VARIANT THEREOI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn version 3.0
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                                                                                                                                                                                                                                                                                                                                                                          / Match 9.1%; Score 12; DB 14; Length 23;
Local Similarity 100.0%; Pred. No. 2.1e+04;
nes 12; Conservative 0; Mismatches 0; Indels
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100.0%; Pred. No. 2.1e+04;
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PRIOR APPLICATION NUMBER: US 60/273,808
PRIOR FILING DATE: 2001-03-07
PRIOR APPLICATION NUMBER: US 60/278,983
PRIOR FILING DATE: 2001-03-27
NUMBER OF SEQ ID NOS: 75
SOFTWARE: Patentin version 3.0
SEQ ID NO 53
LENGTH: 23
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CURRENT FILING DATE: 2002-02-21
PRIOR APPLICATION NUMBER: US 60/270,132
PRIOR FILING DATE: 2001-02-21
PRIOR PRICING DATE: 2001-02-21
PRIOR PRICING DATE: 2001-02-27
PRIOR FILING DATE: 2001-03-27
PRIOR FILING DATE: 2001-03-27
NUMBER OF SEQ ID NOS: 74
SOPTWARE: Patentin version 3.0
SEQ ID NO 43
LENGTH: 23
CURRENT APPLICATION NUMBER: US/10/086,156
CURRENT FILING DATE: 2002-02-28
FPRIOR APPLICATION NUMBER: US 60/272,190
FRIOR FILLING DATE: 2001-02-28
FPRIOR APPLICATION NUMBER: US 60/274,258
FPRIOR FILLING DATE: 2001-03-07
NUMBER OF SEQ ID NOS: 98
SOFTMARE: Patentin version 3.0
SEQ ID NO 73
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US-10-080-980-43
                                                                                                                                                                                                                                                                        Sequence 73, Application US/10086156 Publication No. US20030054989A1 GENERAL INFORMATION:
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Best Local
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TITLE OF INVENTION: POLYNUCLEOTIDE ENCODING TWO NOVEL HUMAN POTASSIUM CHANNEL BETA-S
TITLE OF INVENTION: K+betaM4 and K+betaM5
FILE REFERENCE: D0115NP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Bristol-Myers Squibb Company
TITLE OF INVENTION: POLYNUCLECTIDE ENCODING A NOVEL HUMAN G-PROTEIN COUPLED RECEPTOR
TITLE OF INVENTION: HGBRBMY27
FILE REFERENCE: D0134.NP
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; LENGTH: 23
; TYPE: DMA:
; ORGANISM: Homo sapiens
US-10-086-156-73

Query Match
Best Local Similarity 100.0%; Pred. No. 2.1e+04;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 31 GACTGCGTGAGT 42
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Db 20 GACTGCGTGAGT 9

Search completed: February 3, 2005, 00:11:24

Job time: 75.8037 secs
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      GenCore version 5.1.6 (c) 1993 - 2005 Compugen Ltd
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US-09-171-769A-34
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US-09-866-108A-512
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5112, Ap
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5123, Appl
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127, Appl
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562, Appl
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563, Appl
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US-08-418-848A-9; Sequence 49, Application US/08418848A
; Patent No. 5847096
; Patent No. TRANSPORMATION:
STREET: 345 PARK AVENUE CITY: NEW YORK COUNTRY: U.S.A.
COUNTRY: 0.15.4
ZIP: 10.15.4
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                         APPLICANT: SCHUBERT, MANFRED, HARMISON II,
APPLICANT: GEORGE G., CHANG-JIE, CHEN, BANJERJEA,
TITLE OF INVENTION: DEFECTIVE, INTEFFERING
TITLE OF INVENTION: HIV PARTICLES
                                                                                                                                                NUMBER OF SEQUENCES: 77
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORGAN & FINNEGAN, L.L.P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-705-400-28
US-09-779-5360-9
US-09-79-5360-9
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US-09-198-452A-2711

US-09-198-452A-4232

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US-08-976-161-19
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US-09-517-467B-297
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Sequence 18, Appl
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Sequence 19, Appl
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Sequence 19, Appl
Sequence 10, Appl
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Sequence 18, Appl
Sequence 18, Appl
Sequence 209, Appl
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Sequence 201, Appl
Sequence 3, Appl
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Sequence 11, Appl
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US-08-435-350-67
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                            COMPUTER: IBM COMPATIBLE
COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: IBM P.C. DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/435,350
FILING DATE: 05-MAY-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/936,531
APPLICATION NUMBER: 07/936,531
APPLICATION NUMBER: 07/936,531
APPLICATION NUMBER: 32,327
REGISTRATION NUMBER: 32,327
REGISTRATION NUMBER: 32,327
REGISTRATION NUMBER: 197/245
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEPAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 67, Appli
Patent No. 5599704
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FILING DATE: 07-APR-1995
CLASSIFICATION: 526
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/936,849
FILING DATE: 28-AUG-1992
CLASSIFICATION: 526
ATTORNEY/AGENT INFORMATION:
NAME: RICHARD W. BORK
NAME: RICHARD W. BORK
REGISTRATION NUMBER: 36,459
REFERENCE/DOCKET NUMBER: 2026-409
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: Sir-
TOPOLOGY: 1:
418-848"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: 212-758-4800
TELEPAX: 212-751-6849
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CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: James D. Thompson
APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: METHOD AND REAGENT FOR
TITLE OF INVENTION: TREATMENT OF BREAST CANCER
NUMBER OF SEQUENCES: 118
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 611 West :
CITY: Los Angeles
STATE: California
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Similarity 100.0%;
13; Conservative
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               67-3510
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k; Pred. No. 1.3
0; Mismatches
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; MOLECULE TYPE: other nucleic acid
; DESCRIPTION: /desc = "oligonucleotide"
US-08-702-344-33

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Best Local Similarity 84.6%;
                                                                        Query Match
Best Local Similarity
                                                             Matches
                                                                                                                                                                                            TELEFAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: 33:
SEQUENCE CHARACTERISTICS:
LENGTH: 29 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Merberg, David
APPLICANT: Treacy, Maurice
APPLICANT: Spaulding, Vikki
TITLE OF INVENTION: SECRETED P
TITLE OF INVENTION: ENCODING T
NUMBER OF SEQUENCES: 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   tent No.
                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: Brown, Scott A.
REGISTRATION NUMBER: 32,724
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 498-8224
                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/702,344
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25
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CAGCCCTGCATGC 13
                    CAGCCCTGCATGC 58
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McCoy, John
                                                        Conservative
                                                                                                                                                                 ss: single
linear
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                                                       9.8%; Score 13; DB
100.0%; Pred. No. 1.3
tive 0; Mismatches
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Pred. No. 1.
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                                                                                      DB 1;
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1.3e+03;
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hes 0;
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RESULT 5
US-09-476-387-442
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TITLE OF INVENTION: Nucleotide triphosphate and
FILE REFERENCE: MBHB00-831-B (247/276)
CURRENT APPLICATION NUMBER: US/09/474,432B
CURRENT FILING DATE: 1999-12-19
PRIOR APPLICATION NUMBER: US 60/064,866
PRIOR FILING DATE: 1997-11-05
PRIOR APPLICATION NUMBER: US 60/084,727
PRIOR APPLICATION NUMBER: US 60/084,727
PRIOR FILING DATE: 1998-04-29
PRIOR APPLICATION NUMBER: US 09/186,675
PRIOR APPLICATION NUMBER: US 09/186,675
PRIOR APPLICATION NUMBER: US 09/301,511
PRIOR FILING DATE: 1999-04-28
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                                                                                  PRIOR APPLICATION NUMBER: 09/474, 432
PRIOR FILING DATE: 1999-12-29
PRIOR APPLICATION NUMBER: 09/301,511
PRIOR FILLING DATE: 1999-04-28
PRIOR APPLICATION NUMBER: 09/186,675
PRIOR FILLING DATE: 1998-11-04
PRIOR APPLICATION NUMBER: 60/083,727
PRIOR FILLING DATE: 1998-04-29
PRIOR FILLING DATE: 1998-04-29
PRIOR APPLICATION NUMBER: 60/064,866
PRIOR FILLING DATE: 1997-11-05
            SOFTWARE: Pat
SEQ ID NO 442
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Patent No. 6617438
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SEQ ID NO 443
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Best Local Similarity
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                   APPLICANT: Karpeisky, Alex
APPLICANT: Adamic, Jasenka Matulic
APPLICANT: Sweedler, Dave
APPLICANT: Sweedler, Dave
APPLICANT: Zinnen, Shawn
TITLE OF INVENTION: Nucleotide Triphosphate and their Incorporation into Oligonucleo
FILE REFERENCE: MBHB00-831-C (249/073)
CURRENT APPLICATION NUMBER: US/09/476,387
CURRENT FILING DATE: 2001-04-04
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APPLICANT: Beigelman, Leo
APPLICANT: Burgin, Alex
                                                              NUMBER OF SEQ ID NOS: 1524
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ENGTH: 17
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                                           PatentIn
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Beaudry, Amber
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Adamic, Jasenka
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Beaudry, Amber
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Pred. No.
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; ORGANISM: Homo sapiens 
US-09-476-387-442
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; Sequence 2181, Application US/09866108A
                                                                          US-09-866-108A-2182/c
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CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR PILING DATE: 2000-10-04
PRIOR PILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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Best Local S
Matches 12
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Best Local Similarity
Matches 10; Conserv
              Sequence 2182, Application Patent No. 6686188 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                    Patent No. 6686
SEQ ID NO 2181
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
APPLICANT: GU, Yizhong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: PCT/US01/00663 PRIOR FILING DATE: 2001-01-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: GU, Yizhong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PPLICANT:
                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                     ENGTH:
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12; Conser
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HANZEL, David
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Pred. No. 4.4e+03;
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CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-05-26
PRIOR PPLICATION NUMBER: US 60/236,359
PRIOR PPLICATION NUMBER: PCT/US01/00666
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR PILING DATE: 2001-03-30
PRIOR PILING DATE: 2001-01-30
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APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN
FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR PRIOR DATE: 2000-05-26
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: DT/US01/00666
PRIOR APPLICATION NUMBER: DT/US01/00666
PRIOR APPLICATION NUMBER: DT/US01/00666
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NUMBER OF SEQ ID NOS: 15755
SOFTWARE: Acomica Sequence Listing Engine
Patent No. 6686188
SEQ ID NO 2182
LENGTH: 17
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APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron
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PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART
FILE REFERENCE: AEOMICA-7
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PRIOR APPLICATION NUMBER: PCT/US01/00668
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RANK, David R.
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PENN, Sharron
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100.0%; Pred. No.
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CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR PELION DATE: 2000-05-26
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: DCT/US01/00666
PRIOR FILING DATE: 2001-01-30
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SEQ ID NO 2183
LENGTH: 17
Patent No. 6686
SEQ ID NO 2184
LENGTH: 17
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
                                                              Remaining Prior Application data removed
NUMBER OF SEQ ID NOS: 15755
SOFTWARE: Aeomica Sequence Listing Engine
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PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 100MBER: PCT/US01/00669
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
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PPLICANT: JI, Yonggang
PPLICANT: PENN, Sharron
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ORGANISM: Homo sapiens
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00663
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Patent No. 6686188
SEQ ID NO 2185
LENCTH: 17
TYPE: DNA
ORGANISM: Homo sapiens
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US-09-866-108A-2185/c
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APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, MSIN-LIKE GENE EXPRESSED IN HUMAN HEART .
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART .
FILE REFERENCE: AEOMICA-7
FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108A
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PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
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APPLICATION NUMBER: PCT/US01/00667
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100.0%; Pred. No.
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                                                                                            Length 17;
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                                                              Indels
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; Sequence 2186, Application US/09866108A
; Patent No. 6686188

RESULT 11 US-09-866-108A-2186/c

PPLICANT: GU, Yizhong

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RESULT 12
US-08-379-078-717
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; ORGANISM: Homo sapiens
US-09-866-108A-2186
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CURRENT APPLICATION NUMBER: US/09/866,108A

CURRENT FILING DATE: 2001-05-25

PRIOR APPLICATION NUMBER: US 60/207,456

PRIOR FILING DATE: 2000-05-26

PRIOR FILING DATE: 2000-05-26

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR FILING DATE: 2000-09-27

PRIOR APPLICATION NUMBER: PCT/US01/00666

PRIOR FILING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/US01/00667

PRIOR APPLICATION NUMBER: PCT/US01/00667

PRIOR APPLICATION NUMBER: PCT/US01/00664

PRIOR APPLICATION NUMBER: PCT/US01/00669

PRIOR APPLICATION NUMBER: PCT/US01/00669

PRIOR APPLICATION NUMBER: PCT/US01/00669

PRIOR APPLICATION NUMBER: PCT/US01/00665

PRIOR FILING DATE: 2001-01-30

PRIOR FILING DATE: 2001-01-30

PRIOR FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Patent No. 66801
SEQ ID NO 2186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 100.
12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Patent No. 5639612
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 717, Appl Patent No. 5639612
                   ZIP: 92660

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 15755
SOFTWARE: Acomica Sequence Listing Engine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Mitsuhashi, Masato
APPLICANT: Cooper, Allan
TITLE OF INVENTION: Gene Detection System
NUMBER OF SEQUENCES: 726
                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 726
CORRESPONDENCE ADDRESS:
ADDRESSEE: KNOBBE, MARTENS, OLSON AND BEAR
STREET: 620 Newport Center Drive 16th Floor
APPLICATION NUMBER: US/08/379,078
                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                             CITY: Newport Beach
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PENN, Sharron G.
HANZEL, David K.
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US-08-860-882A-52
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                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy Disk
COMPUTER: IBM compatible
OPERATING SYSTEM: DC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/860,882A
FILING DATE: JUNE 23, 1997
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: DONALD J. BIRD
REGISTRATION NUMBER: 25,323
REFERENCE/DOCKET NUMBER: 9901/238653
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 861-3027
TETLEPHONE: (202) 861-3027
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No.
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Best Local Similarity
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TELEPHONE: 714-760-0404
TELEPAX: 714-760-9502
INFORMATION FOR SEQ ID NO: 717:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: DAVIES, I TITLE OF INVENTION: NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 12-NOV-1992
ATTORNEY/AGENT INFORMATION:
NAME: Altman, Daniel E.
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
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           TELEPHONE: (202)
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                                                                                                                                                                                                                                                                            STATE: D
                                                                                                                                                                                                                                                                                                                         STREET:
                                                                                                                                                                                                                                                                                                            CITY: WASHINGTON
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REGISTRATION NUMBER: 34
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                         ADDRESSEE:
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10. 59852
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USA
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: (202) 822-0944
6174627 CUSH
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MARSHAM, PETER ROBERT
HEATON, DAVID WILLIAM
DAVIES, DAVID HUW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TAYLORSON, CHRISTOPHER JOHN EGGELTE, HENDRIKUS JOHANNES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                HENNAM, JOHN FREDERICK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FRANCIS THOMAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BRIAN ROBERT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      9.1%; Score 12;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                         BURY, MADISON & SUTRO YORK AVENUE, N.W.
                                                                                                                                                                                                                                                                                                                                                                                     CHEMICAL COMPOUNDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   S
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. 4.4e+03;
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Sequence 289, Application US/07974409C
Patent No. 6300058
GENERAL INFORMATION:
APPLICANT: Akitaya, Tatsuo
APPLICANT: Mitsuhashi, Massato
APPLICANT: Cooper, Alla
ITITLE OF INVENTION: METHOD AND REAGEN
TITLE OF INVENTION: FOR MEASURING MES
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                                                                                                                                                                    US-07-974-409C-289
                                                                                                                                                                              RESULT 15
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US-09-171-945-110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-860-882A-52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 9.1%;
Best Local Similarity 100.0%;
Matches 12; Conservative (
                                                                                                                                                                                                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 19 bases
                                                            NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: singl
TOPOLOGY: linear
                          STREET: 620 Newport CITY: Newport Beach
                                                                                                                                                                                                                                                                  Local
COUNTRY: U
                                                   ADDRESSEE:
                                                                                                                                                                                                                                  18 GGACCTGCTGCA 29
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                                                                                                                                                                                                                                                       l Similarity
12; Conserv
  . CA
USA
                                                                                                                                                                                                              GGACCTGCTGCA 12
                                        E: Knobbe, Martens, Olson, and Bear 620 Newport Center Dr. Sixteenth Floor
                                                                                                                                                                                                                                                       Conservative
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                                                                                 METHOD AND REAGENT
FOR MEASURING MESSENGER RNA
                                                                                                                                                                                                                                                     9.1%; Score 12; DB 3; L
100.0%; Pred. No. 4.4e+03;
tive 0; Mismatches 0;
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; Pred. No.
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REGISTRATION NUMBER: 34,115
REFERENCE DOCKET NUMBER: HITA
TELECOMMUNICATION INFORMATION:
TELEPHONE: 714-760-0404
TELEPAX: 714-760-9502
INFORMATION FOR SEQ ID NO: 289:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 34, Applicati
Patent No. 6436691
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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FILING DATE: 12-NOV-1992
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: // IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
           COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: MS WOR'D
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/011,769A
FILING DATE: 13-Feb-1998
CLASSIFICATION LOTA:
APPLICATION DATA:
APPLICATION NUMBER: PCT/GB96/01975
FILING DATE: 13-NUG-1996
APPLICATION NUMBER: GB 9612295.7
FILING DATE: 12-UN-1996
APPLICATION NUMBER: GB 9611019.2
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TOPOLOGY: lin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH:
                                                                                                                                                                                                                                                         ZIP: 20005
COMPUTER READABLE FORM:
MEDIUM TYPE: 1.44 /
                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Chemical Compounds NUMBER OF SEQUENCES: 87
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Similarity 100.0%; Pred. No. 4.4e+
                                                                                                                                                                                                                                                                                                               CITY: Washington
STATE: D.C.
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nucleic acid
                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Pillsbury Madison & Sutro, LLP STREET: 1100 New York Ave., N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Application US/09011769A
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HENNAM, John F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HENNEQUIN, Laurent F.A. MARSHAM, Peter R. DOWELL, Robert I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cDNA to mRNA
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                                                                                                                                                                                                                                                             1.44 Mb disk
25-MAY-1996
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FILING DATE:

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RESULT 18
US-08-930-001-6/c
. sequence 6, Application US/08930001A
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; ANTI-SENSE: NO
PCT-US93-00977-289
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PCT-US93-00977-289
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                                                                                                                                                                                            Query Match
Best Local (
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GENERAL INFORMATION:
TITLE OF INVENTION: METHOD AND REAGENT FOR MEASURING MESSENGER RNA
NUMBER OF SEQUENCES: 711
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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TELEFAX: 714-760-9502
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 34:
                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Altman, Daniel E.
REGISTRATION NUMBER: 34,115
REFERENCE/DOCKET NUMBER: HITACHI.006H
TELECOMMUNICATION INFORMATION:
TELEPHONE: 714-760-0404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/00977
FILING DATE: 19930129
CLASSIFICATION:
                                                                                                                                                                                                                                                                                         MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Knobbe, Martens, Ol:
STREET: 620 Newport Center Dr.
                                                                                                                                                                                                                                                                                                          TOPOLOGY:
                                                                                                                                                                                                                                                                                                                         STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                        TYPE: NUCLEIC ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Match 9.1%; So Local Similarity 100.0%; Fig. 12; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Newport Beach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                                                                                                                                                                       ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear MOLECULE TYPE: other nucleic acid SEQUENCE DESCRIPTION: SEQ ID NO: 34:
                                                                                                                                           15 CCAGGACCTGCT 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18 GGACCTGCTGCA 29
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                                                                                                                                                                          l Similarity
12; Conserv
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TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                               CCAGGACCTGCT 17
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Conservative (
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                                                                                                                                                                                                                                                                                           cDNA to mRNA
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Pred. No.
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Dr. Sixteenth Floor
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No. 4.4e+03;
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. 4.4e+03;
ches 0;
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FILE REFERENCE: 0230-118P
CURRENT APPLICATION NUMBER: US/08/930,001A
CURRENT FILING DATE: 1997-09-26
EARLIER APPLICATION NUMBER: JAPAN 106819/1995
EARLIER FILING DATE: 1995-03-27
NUMBER OF SEQ ID NOS: 8
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 6
LENGTH: 20
                                                                                                                                      RESULT 20
US-09-920-759-53/c
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US-09-640-101-99
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                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: P.
SEQ ID NO 99
LENGTH: 20
TYPE: DNA
Sequence 53, Application US/09920759
Patent No. 6537811
GENERAL INFORMATION:
APPLICANT: Brenda F. Baker
APPLICANT: Susan M. Freier
TITLE OF INVENTION: ANTISENSE MODULATION OF SAP-1 EXPRESSION
FILE REFERENCE: RTS-0267
CURRENT APPLICATION NUMBER: US/09/920,759
                                                                                                                                                                                                                                                                                         Query Match
Best Local 9
                                                                                                                                                                                                                                                                          Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Antisense Modulation of p38 Mitogen FITLE OF INVENTION: Activated Protein Kinase Expression FILE REFERENCE: ISPH-0488
CURRENT APPLICATION NUMBER: US/09/640,101
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: 09/286,904
PRIOR FILING DATE: 1999-04-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: MURATA, NO. 628141210
TITLE OF INVENTION: METHOD FOR PRODUCING OSMOTOLERANT PLANTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Description of Artificial Sequence:primer
                                                                                                                                                                                                                                                                                         Local Similarity
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Nero, Pamela S.
McKay, Robert
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/ 100.0%; Pr
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; Pred. No.
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Pred. No.
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                                                                                                                                                                                                                                                                                                         DB 4;
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S
                                                                                                                                              ; TYPE: DNA
; ORGANISM: Chlamydia pneumoniae
US-09-198-452A-5261
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; ORGANISM: Chlamydia pneumoniae
US-09-198-452A-5153
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US-09-198-452A-5153
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US-09-920-759-53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Griffels, R.

APPLICANT: Griffels, R.

TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence and polypeptides, fragments
TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnosis, preve
TITLE OF INVENTION: and treatment of infection
FILE REFERENCE: 9710-003-999
CURRENT APPLICATION NUMBER: US/09/198,452A
CURRENT APPLICATION NUMBER: US/09/198,452A
CURRENT FILING DATE: 1998-11-24
NUMBER OF SEQ ID NOS: 6849
SEQ ID NO 5153
LENGTH: 20
                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/198,452A
CURRENT FILING DATE: 1998-11-24
NUMBER OF SEQ ID NOS: 6849
SEQ ID NO 5261
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                Sequence 5261, Applic Patent No. 6559294 GENERAL INFORMATION:
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NUMBER OF SEQ ID NOS:
SEQ ID NO 53
LENGTH: 20
                                                                         Matches
                                                                                           Query Match
Best Local
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                                                                                                                                                                                                                                                                                                 APPLICANT: Griffais, R. TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence and polypeptides, fragments TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnosis, preveTITLE OF INVENTION: and treatment of infection
FILE REFERENCE: 9710-003-999
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18
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12; Conserv
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12; Conserv
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GCATCATCTGCC 7
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                                 GCATCATCTGCC 16
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larity 100.0%; Pred. No. 4.4e+03;
Conservative 0; Mismatches 0;
                                                                       Conservative
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                                                                   9.1%; Score 12; DB 4; Lo
100.0%; Pred. No. 4.4e+03;
tive 0; Mismatches 0;
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. 4.4e+03;
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RESULT 25
US-09-329-515A-59
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US-09-724-126A-29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; Sequence 6, Application US/09091885; Patent No. 6756525; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-091-885-6/c
                                                                                             Sequence 59, Application US/09329515A Patent No. 6740487 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 12; Conserv
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LENGTH: 21
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APPLICANT: Schwartz, David A.
APPLICANT: Schutte, Brian C.
APPLICANT: Schutte, Brian C.
TITLE OF INVENTION: Variant TLR4 nucleic acid and uses thereof
FILE REFERENCE: 875.0100031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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Best Local :
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APPLICANT: Kwak, Keith
TITLE OF INVENTION: Human E3 Alpha Ubiquitin Ligase Family
FILE REFERENCE: 01017/35966A
CURRENT APPLICATION NUMBER: US/09/724,126A
CURRENT FILING DATE: 2000-11-28
PRIOR APPLICATION NUMBER: US 60/187,211
PRIOR APPLICATION NUMBER: US 60/187,211
PRIOR FILING DATE: 1999-03-01
NUMBER OF SEQ ID NOS: 29
SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: MURATA, NO. 675652510
TITLE OF INVENTION: METHOD FOR PRODUCING TEMPERATURE-TOLERANT PLANTS
FILE REFERENCE: 230-123P
CURRENT APPLICATION NUMBER: US/09/091,885
CURRENT FILING DATE: 1998-06-26
NUMBER OF SEQ ID NOS: 8
NUMBER OF SEQ ID NOS: 8
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Artificial sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION:
                                                                                                                                                                                                                                                                                49
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Similarity 100.0%; Pred. No.
12; Conservative 0; Mismatcl
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100.0%; Pred. No.
tive 0; Mismatch
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hes 0;
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Best Local Similarity
Matches 12; Conserve
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; ORGANISM: Homo sapiens
US-09-329-515A-59
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                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 27
US-08-211-202-76/c
                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: HOGENBOOM; Hendricus Renerus Jacc
APPLICANT: BAIER, Michael
APPLICANT: JESPERS, Laurent Stephane Anne The
APPLICANT: WINTER, Gregory Paul
APPLICANT: WINTER, Gregory Paul
TITLE OF INVENTION: Production of chimeric &
TITLE OF INVENTION: combinatorial approach
NUMBER OF SEQUENCES: 144
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-339-775-4
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LENGTH: 22
TYPE: DNA
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APPLICANT: Lex M. COWBERT
TITLE OF INVENTION: ANTISENSE MODULATION OF G-ALPHA-I3 EXPRESSION
FILE REFERENCE: RTS-0069
CURRENT APPLICATION NUMBER: US/09/339,775
CURRENT FILING DATE: 1999-06-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 59
LENGTH: 21
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Patent No. 6063626
                                                                                                                                                                                                                                                                                                                                                                                                             Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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ZIP: 60606-6402
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/211,202
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
OTHER INFORMATION: PCR Probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
                                                                                                                                                              CITY: Chicago
STATE: Illino
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                                                                                                                                                                                                                                   ADDRESSEE: David W. Clough, Marshall O'Toole Gerstein Murray &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    45 GCAGCCCTGCAT 56
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                                                                                                                                                             Illinois
                                                                                                                                                                                                6300 Sears Tower,
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Conservative
                                                                                                                                              USA
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100.0%; Pred. No. 4.4e+03;
                                                                                                                                                                                                                                                                                                                                        Laurent Stephane Anne Therese
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Pred. No. 4.4e+03;
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                                                                                                                                                                                                233 South Wacker Drive
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                                                                                                                                                                                                                                                                                                         chimeric antibodies
                                      Version
                                      #1.25 (EPO)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
              ZIP: 60606-6402
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPUTER:
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: PATENTIN Release #1.0, V.
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: PCT/GE
FILING DATE: 15-MAY-1992
ATTORNEY/AGENT INFORMATION:
NAME: David W. Clough
REGISTRATION NUMBER: 36,10
REFERENCE/DOCKET NUMBER: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 24-MAR-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9:
FILING DATE: 24-MAR-1992
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 23-SEP-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 91
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: David W. Clough
STREET: Marshall, O'Toole,
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APPLICANT:
APPLICANT:
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APPLICANT:
APPLICANT:
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TELEPHONE: 312-474-6300
TELEPAX: 312-474-0448
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APPLICATION NUMBER: GB 9206318.9
                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Methods for producing members of specific princeNTION: binding pairs
                                                                                                                                                                                           ADDRESSEE:
STREET: Ma:
STREET: 63
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                                                                                                                                                            STATE:
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                                                                                                                                       COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 554, Application US/08350260A
o. 5962255
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12; Conservative
                                                                                                                                                          Chicago
Illinois
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                                                                                                                                                                                           Marshall, O'Toole, Gerstein, Murray & Borun
6300 Sears Tower, 233 South Wacker Drive
                                                                                                                                                                                                                                                                                                                  Johnson, Kevin Stuart
Smith, Andrew John Hammond
                                                                                                                                           USA
                                                                                                                                                                                                                                                                                                                                                                                         Griffiths, Andrew David Williams, Samuel Cameron
                                                                                                                                                                                                                                                                                                                                                       Nissim, Ahuva
                                                                                                                                                                                                                                                                                                                                                                            Waterhouse, Peter
                                                                                                                                                                                                                                                                                                                                                                                                                                  Winter,
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                                                                                                                                                                                                                                                                                                                                                                                                             Gregory Paul
hs, Andrew David
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100.0%; Pred. No.
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US/08/350,260A
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                                     Version #1.25 (EPO)
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FILING DATE: 15-MAY-

15-MAY-1991

GB 9110549.4

PRIOR APPLICATION DATA:

APPLICATION NUMBER:

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US-08-860-882A-39/c
; Sequence 39, Application US/08860882A
; Patent No. 5985281
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APPLICANT: TAYLORS:
APPLICANT: EGGELTE
APPLICANT: TARRAGO:
APPLICANT: RABIN, I
APPLICANT: BOYLE, I
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                                                                                                                                                     APPLICANT: BLAKELY, DAVID CHARLES
APPLICANT: MARSHAM, PETER ROBERT
APPLICANT: HEATON, DAVID WILLIAM
APPLICANT: DAVIES, DAVID HUW
TITLE OF INVENTION: CHEMICAL COMPOUN
NUMBER OF SEQUENCES: 77
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: 312-474-6300 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 08/307,619 FILING DATE: 16-SEP-1994 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US 01 FILING DATE: 31-MAR-1994 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: PCT/GB92/00883
FILING DATE: 15-MAY-1992
PRIOR APPLICATION DATA:
APPLICATION UNMBER: PCT/GB93/00605
FILING DATE: 24-MAR-1993
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy Disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 10-JUL-
PRIOR APPLICATION DATA:
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                                                                                                                      ADDRESSEE:
STREET: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Clough, David W
REGISTRATION NUMBER: 36,107
REFERENCE/DOCKET NUMBER: 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
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                                                      ZIP: 20005
                                                                       COUNTRY:
                                                                                                        CITY: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ENGTH:
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                                                                                                                        1100 NEW YORK AVENUE, N.W.
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                                                                                                                                                                                                                                                                                RABIN, BRIAN ROBERT
BOYLE, FRANCIS THOMAS
HENNAM, JOHN FREDERICK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                         USA
                                                                                                                                                                                                                                                                                                                                                    TAYLORSON, CHRISTOPHER JOHN EGGELTE, HENDRIKUS JOHANNES
                                                                                                                                                                                                                                                                                                                                     TARRAGONA-FIOL, ANTONIO
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IBM compatible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       9.1%; Score 12; DB 2;
100.0%; Pred. No. 4.4e+03
                                                                                                                                                                                              CHEMICAL COMPOUNDS
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RESULT 30
US-08-860-882A-51/c
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Best Local Similarity 100.0%; P
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TELEPAX: (202) 822-0944.
TELEX: 6174627 CUSH
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                             ATTORNEY/AGENT INFORMATION:

NAME: DONALD J. BIRD
REGISTRATION UMBER: 25,323
REFERENCE/DOCKET NUMBER: 9901
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 861-3027
TELEPHONE: (202) 861-3027
                                                                                                                                OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/860,882A
FILING DATE: JUNE 23, 1997
CLASSIFICATION: 424
CLASSIFICATION: 424
CLASSIFICATION: 424
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APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy Disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF
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                                                                                                                                                                                                                               COMPUTER: IBM compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                        STREET: 1100 NEW CITY: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: singl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: DONALD J. BIRD REGISTRATION NUMBER: 25, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM: PC-DOS/MS-DOS
TELEFAX: (202) 822-0944
TELEX: 6174627 CUSH
                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                         STATE: D.C
                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE:
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1100 NEW YORK AVENUE, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RABIN, BRIAN ROBERT
BOYLE, FRANCIS THOMAS
HENNAM, JOHN FREDERICK
BLAKELY, DAVID CHARLES
MARSHAM, PETER ROBERT
                                                                                                                                                                                                                                                                                                                          USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TAYLORSON, CHRISTOPHER JOHN EGGELTE, HENDRIKUS JOHANNES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TARRAGONA-FIOL, ANTONIO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
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100.0%; Pred. No.
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                                                                        9901/238653
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; TOPOLOGY: linear; SEQUENCE TYPE: other nucleic acid; SEQUENCE DESCRIPTION: SEQ ID NO: 6: US-09-011-769A-6
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US-09-011-769A-6/c
; Sequence 6, Application US/09011769A
; Patent No. 6436691
; GENERAL INFORMATION:
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                                                               Query Match
Best Local S
Matches 12
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                                                                                                                                                                                                                                                   APPLICATION NUMBER: PCT/GB96/01975
FILING DATE: 13-AUG-1996
APPLICATION NUMBER: GB 9612295.7
FILING DATE: 12-JUN-1996
APPLICATION NUMBER: GB 9611019.2
FILING DATE: 25-MAY-1996
APPLICATION NUMBER: GB 9516810.0
FILING DATE: 16-AUG-1995
FORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: singl
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ENGTH:
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MEDLIM TYPE: 1.44 Mb disk

COMPUTER: IEM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: MS WORD

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/09/011,769A
FILING DATE: 13-Feb-1998
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Pillsbury Madison & Sutro, LLP
STREET: 1100 New York Ave., N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MARSHAM, Peter R.
DOWELL, Robert I.
TITLE OF INVENTION: Chemical Compounds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
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12; Conserv
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                             GGACCTGCTGCA 29
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TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: U.S.A.
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                                                                 Conservative
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HENNAM, John F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20005
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100.0%; Pred. No. 4.4e+03;
                                                                               100.0%;
                                                               9.1%; Score 12; DB 4; L
100.0%; Pred. No. 4.4e+03;
ive 0; Mismatches 0;
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RESULT 32
US-09-011-769A-33/c
US-09-011-769A-33/c
; Sequence 33, Application
; Patent No. 6436691
; Patent INFORMATION:
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US-09-104-337A-554/c
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                                                                                                                                                                     Sequence 554, Application US/09104337A Patent No. 6492160 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 16-AUG-1995 INFORMATION FOR SEQ ID NO: 33:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:

APPLICATION NUMBER: PCT/GB96/01975

FILING DATE: 13-AUG-1996

APPLICATION NUMBER: GB 9612295.7

FILING DATE: 12-UN-1996

APPLICATION NUMBER: GB 9611019.2

FILING DATE: 25-MAY-1996

APPLICATION NUMBER: GB 9516810.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear MOLECULE TYPE: other nucleic acid SEQUENCE DESCRIPTION: SEQ ID NO: 33:
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MEDIUM TYPE: 1.44 mb disk
COMPUTER: IEM PC compartible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: MS WOR'D
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Chemical Compounds NUMBER OF SEQUENCES: 87
CORRESPONDENCE ADDRESS:
                         TITLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                      APPLICANT: Winter,
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                                                                                                                                                                                                                                                                                                                                                                                12;
                                                                                                                                                                                                                                                                                                                                           18 GGACCTGCTGCA 29
                                                                                                                                                                                                                                                                                                      19 GGACCTGCTGCA 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 23 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/09/011,769A
FILING DATE: 13-Feb-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 20005
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                       INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HENNEQUIN, Lau-
                       Johnson, Kevin Students Smith, Andrew John Hammond
                                                                                                             Winter, Gregory Paul
Griffiths, Andrew David
Williams, Samuel Cameron
                                                                             Waterhouse, Peter
Nissim, Ahuva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DAVIES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MARSHAM, Peter
DOWELL, Robert
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100.0%; Pred. No. 4.4e+03
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Methods for producing members of specific binding pairs
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RESULT 34
US-10-067-443-48/c
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NAME: Bartnicki, Audrey L.

REGISTRATION NUMBER: 40,499

REFERENCE/DOCKET NUMBER: 28111/32372A

TELECOMMUNICATION INFORMATION:

TELEPHONE: 312-474-6300

INFORMATION FOR SEQ ID NO: 554:

SEQUENCE CHARACTERISTICS:

LENGTH: 23 base pairs

TYPE: nucleic acid

STRANDEDNESS: single
GENERAL INFORMATION:
APPLICANT: Bristol-Myers Squibb Company
ITTLE OF INVENTION: POLYNUCLECTIDE ENCODING A NOVEL METALOPROTEASE HIGHLY EXPRESSED
TITLE OF INVENTION: SPINAL CORD, MP-1
FILE REFERENCE: D0073 NP
CURRENT FILING DATE: 2002-02-05
PRIOR APPLICATION NUMBER: US/10/067,443
CURRENT FILING DATE: 2001-02-05
PRIOR FILING DATE: 2001-02-05
PRIOR FILING DATE: 2001-02-05
PRIOR FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 71
SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                                                                                          Sequence 48, Application US/10067443 Patent No. 6642041
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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APPLICATION NUMBER: US/09/104,337A
FILING DATE: 25-un-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 600
CORRESPONDENCE ADDRESS:
ADDRESSEE: Madrey L. Bartnicki
STREET: Marshall, Gerstein & Borun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE DESCRIPTION: SEQ ID NO: 554:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA
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                                                                                                                                                                                                                                                                                                                                                                                                                                         31 GACTGCGTGAGT 42
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 08/350,260 FILING DATE: 05-DEC-1994 APPLICATION NUMBER: GB 9110549.4 FILING DATE: 15-MAY-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: Illinois
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 16-SEP-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: GB 9206318.9
FILING DATE: 24-MAR-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 31-MAR-1994
APPLICATION NUMBER: US 08/307,619
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 15-MAY-1992
APPLICATION NUMBER: PCT/GB93/00605
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/ 100.0%; Pr
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RESULT 36
US-09-724-126A-28
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US-10-153-064-48/c
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LENGTH: 23
TYPE: DNA """
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 60/293,212
PRIOR FILING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 137
SOFTWARE: Patentin version 3.1
SEQ ID NO 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Bell et al.
TITLE OF INVENTION: Chemokine Beta-1 Fusi-
FILE REPERENCE: PF556
CURRENT APPLICATION NUMBER: US/10/153,064
CURRENT FILING DATE: 2002-05-24
Query Match 9.1%; Score 12;
Best Local Similarity 100.0%; Pred. No.
Matches 12; Conservative 0; Mismatcl
                                                                                                                                                                    SOFTWARE: PatentIn version 3.0
SEQ ID NO 28
LENGTH: 23
                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 28, Application US/09724126A Patent No. 6706505
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Best Local Similarity
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APPLICANT: Kwak, Keith
TITLE OF INVENTION: Human E3 Alpha Ubiquitin Ligase Family
FILE REFERENCE: 01017/35966A
CURRENT APPLICATION NUMBER: US/09/724,126A
CURRENT FILLNG DATE: 2000-11-28
PRIOR APPLICATION NUMBER: US 60/187,211
PRIOR APPLICATION NUMBER: US 60/187,211
PRIOR FILING DATE: 1999-03-01
NUMBER OF SEQ ID NOS: 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 23
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Degenerate Vkappa forward primer
OTHER INFORMATION: amplifying human VI domains
                                                                                      OTHER INFORMATION: Description of Artificial Sequence: PCR Primer
                                                                                                                FEATURE:
                                                                                                                                  ORGANISM: Artificial sequence
                                                                                                                                                      TYPE: DNA
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Local Similarity 100.0%; Pred. No. 4.4e+03;
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100.0%; F1
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100.0%; Pred. No. 4.4e+03
ive 0; Mismatches 0
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                                     DB 4;
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                 4.4e+03;
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RESULT 37 US-09-123-030-33

Sequence 33,

Application US/09123030

Patent No. 6365337 GENERAL INFORMATION:

APPLICANT: Lette, Verity A. APPLICANT: Frankel, Mayne N. APPLICANT: Campbell, Kevin P. APPLICANT: Fellx, Ricardo APPLICANT: Biddlecome, Gloria

TITLE OF INVENTION: Genes Encoding Neuronal Voltage-Gated Calcium Channel TITLE OF INVENTION: Gamma Subunits FILE REFERENCE: US App. 09/123,030 CURRENT APPLICATION UNMERR: US/09/123,030 CURRENT FILING DATE: 1998-07-27 NUMBER OF SEQ ID NOS: 40 SOFTWARE: Patentin Ver. 2.0 SEQ ID NO 3 3 LENGTH: 24

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                                                             PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR PPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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Best Local Similarity
Matches 12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
APPLICANT: HANZEL, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REPERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: COND-05-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Description of Artificial Sequence: primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
                                        APPLICATION NUMBER: PCT/US01/00665
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NUMBER: PCT/US01/00668
                          2001-01-30
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100.0%; Pred. No. 4.
tive 0; Mismatches
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Patent No. 6686188
SEQ ID NO 5110
LENGTH: 25
TYPE: DNA
ORGANISM: Homo sapiens
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN
FILE REFERENCE: ABONICA-7
CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: DCT/US01/00666
PRIOR APPLICATION NUMBER: DCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: DCT/US01/00667
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PRIOR APPLICATION NUMBER: DCT/US01/00667
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                                                                              ; ORGANISM: Homo sapiens US-09-866-108A-5111
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                                                                                                                     ; Patent No. 6686188
; SEQ ID NO 5111
; LENGTH: 25
; TYPE: DNA
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APPLICANT:
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Matches
                 Query Match
Best Local :
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PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
                                                                                                                                                                                                                          Remaining Prior Application data removed NUMBER OF SEQ ID NOS: 15755
                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
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                                                                                                                                                                                                    SOFTWARE: Acomica Sequence Listing Engine
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APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
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11 Similarity
12; Conserv
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Conservative
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N, Sharron
                   9.1%;
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100.0%; Pred. No. 4.
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                 Score 12;
Pred. No.
  Mismatches
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                   DB 4; 1
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PRIOR APPLICATION NUMBER: US 60/207,456

PRIOR APPLICATION NUMBER: US 60/207,456

PRIOR PRILING DATE: 2000-05-26

PRIOR FILING DATE: 2000-10-04

PRIOR PRILING DATE: 2000-10-04

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR APPLICATION NUMBER: CTT/US01/00666

PRIOR APPLICATION NUMBER: PCT/US01/00667

PRIOR PRILING DATE: 2001-01-30

PRIOR PRILING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/US01/00664

PRIOR APPLICATION NUMBER: PCT/US01/00664

PRIOR APPLICATION NUMBER: PCT/US01/00664
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US-09-866-108A-5113/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108A-5112
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; SEQ ID NO 5112
                                                                                                                                                                        Sequence 5113, Application US/09866108A Patent No. 6686188 GENERAL INFORMATION:
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Best Local
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                                                                                         APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggan
APPLICANT: PENN, Shar:
APPLICANT: HANZEL, Dav
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ITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
ILE REFERENCE: AEOMICA-7
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SHANNON, Mark
INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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                                              JI, Yonggang
PENN, Sharron G.
HANZEL, David K.
RANK, David R.
CHEN, Wensheng
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HANZEL, David K.
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Conservative
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APPLICANT: HANZEL, David K.
APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, MAYK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN
FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108A-5113
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SEQ ID NO 5113
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PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING NUMBER: US 60/236,359
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PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
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APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron
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NUMBER OF SEQ ID
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                                                             APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00663
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APPLICATION NUMBER: PCT/US01/00669
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SEQ ID NOS: 15755
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NUMBER: PCT/US01/00665
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108A-5115
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   Query Match
Best Local Similarity
Matches 12; Conserv
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SEQ ID NO 5115
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Patent No. 6686188
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Best Local Similarity
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PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
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NUMBER OF
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
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PRIOR FILING DATE: 2001-01-30
                                                                                                                                 LENGTH: 25
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APPLICATION NUMBER: PCT/US01/00664
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Pred. No.
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       Mismatches
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US-09-866-108A-5117/c
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SEQ ID NO 5116
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PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR PILING DATE: 2000-10-04
PRIOR PILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
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APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
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TYPE: DNA
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CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: PCT/US01/00668
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APPLICATION NUMBER: PCT/US01/00669
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I: SHANNON, Mark
INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
                                                                                                                                                                                                                                                                                                                                                                           15 CCAGGACCTGCT 26
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                                                  PENN, Sharron G
HANZEL, David K
RANK, David R.
CHEN, Wensheng
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100.0%; Pred. No.
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CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25

FILE REFERENCE: AEOMICA-7

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                                                                          CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR PRIOR PRIOR DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
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US-09-866-108A-5118/c
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PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-10-04
PRIOR PILING DATE: 2000-09-27
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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APPLICANT: JI, Yonggan
APPLICANT: PENN, Sharr
APPLICANT: HANZEL, DAV
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Best Local
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SEQ ID NO 5117
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN
FILE REFERENCE: AEOMICA-7
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TYPE: DNA
ORGANISM: Homo sapiens
                               APPLICATION NUMBER: PCT/US01/00669 FILING DATE: 2001-01-30
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1 Similarity 100.08;
12; Conservation
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HANZEL, David K.
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PCT/US01/00665
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Pred. No
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SOFTWARE: Aeomica Sequence, Patent No. 6686188; SEQ ID NO 5119; LENGTH: 25; TYPE: DNA CRGANISM: Homo sapiens
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US-09-866-108A-5119/c
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Query Match 9.1%; Sometime 100.0%; I Best Local Similarity 100.0%; I Matches 12; Conservative 0;
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Patent No. 6686188
SEQ ID NO 5118
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
                                                                                                                                                                                                                         Remaining Prior Application NUMBER OF SEQ ID NOS: 15755
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APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
FILE REFERENCE: AEOMICA-7
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APPLICANT:
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Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 15755
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PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING;DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
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TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                               APPLICATION NUMBER: PCT/US01/00663 FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00667
FILING DATE: 2001-01-30
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5. 6686188
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PENN, Sharron
HANZEL, David
                                                                                                                                                                                                                                              Application data removed -
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                     Score 12;
Pred. No.
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                                     Length 25;
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US-09-866-108A-5120/c

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RESULT 48

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RESULT 49
US-09-866-108A-5121/c
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PRIOR APPLICATION NUMBER: US 60
PRIOR APPLICATION NUMBER: PCT/US
PRIOR APPLICATION NUMBER: PCT/US
PRIOR FILLING DATE: 2001-01-30
PRIOR FILLING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US
PRIOR FILLING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 15755
SOFTWARE: Accomica Sequence Listing Engine
Patent No. 6686188
SEQ ID NO 5120
LENGTH: 25
                                                                                                                                               Patent No. 666011
Patent No. 666011
PRINTERAL INFORMATION:
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Best Local
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APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
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APPLICANT: JI, Yongga
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PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OR APPLICATION NUMBER: PCT/US01/00666
OR FILING DATE: 2001-01-30
OR APPLICATION UNMBER: PCT/US01/00667
OR FILING DATE: 2001-01-30
OR APPLICATION UNMBER: PCT/US01/00664
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00669
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RANK, David R. CHEN, Wensheng SHANNON, Mark
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PENN, Sharron G.
HANZEL, David K.
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US-09-866-108A-5122/c
                                                               CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR PRILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR PILING DATE: 2000-09-27
PRIOR PILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: DCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: DCT/US01/00667
PRIOR APPLICATION NUMBER: DCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Aeomica Sequence Listing Engine
Patent No. 6686188
SEQ ID NO 5121
LENGTH: 25
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 5122, Application US/09866108A Patent No. 6686188
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                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
FILE REFERENCE: AEOMICA-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: GU, Yizhong APPLICANT: JI, Yongga
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
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PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity 100.0%;
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APPLICATION NUMBER: PCT/US01/00667
                                               APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
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PENN, Sharron G
HANZEL, David K
RANK, David R.
CHEN, Wensheng
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L00.0%; Pred. No. 4.4e+03;
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APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN
FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR PRICIATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: GB 024263.6
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: DCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: DCT/US01/00667
PRIOR APPLICATION NUMBER: DCT/US01/00667
PRIOR FILING DATE: 2001-01-30
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; ORGANISM: Homo sapiens
US-09-866-108A-5122
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Patent No. 6686188
SEQ ID NO 5122
LENGTH: 25
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Best Local Similarity
                            Query Match
                                                                       ORGANISM: Homo sapiens -09-866-108A-5123
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PRIOR TILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
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PRIOR FILING DATE: 2001-01-30
Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 15755
SOFTWARE: Acomica Sequence Listing Engine
                                                                                                                                                                                                      Remaining Prior Application data removed - NUMBER OF SEQ ID NOS: 15755
SOFTWARE: Acomica Sequence Listing Engine
                                                                                                                                                                                                                                                                                            PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
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                                                                                                                    LENGTH: 25
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>. 6686188
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HANZEL, David K.
RANK, David R.
CHEN, Wensheng
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100.0%; Pred. No.
  100.0%;
  9.1%; Score 12;
100.0%; Pred. No.
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  DB 4; Length 25; 4.4e+03;
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RESULT 53
US-08-486-857-3/c
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US-08-859-998-358/c
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                                                                                                                                                                    ; MOLECULE TYPE: DNA
; FEATURE:
; OTHER INFORMATION: oligonucleotide primer
US-08-859-998-358
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                                                                                                                            Query Match
Best Local :
                                                                                                               Matches
                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Field, Bree E.
REGISTRATION NUMBER: 37,620
REFERENCE/DOCKET NUMBER: 0909
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-322-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 21-MAY-19
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA: APPLICATION NUMBER: US
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Chenchik, Alex
APPLICANT: Jokhadze, George
APPLICANT: Bibilashvilli, Ro
                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM CO
OPERATING SYSTEM:
                                                                                                                                                                                                                            TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE:
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ZIP: 94025
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Menlo Park
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                                                                            109 TGTCCTACCTGC 120
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                                                                                                                            Similarity
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2200 Sand Hill Road,
                                                                                                                                                                                                                                                                   26 base pairs
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                                                                                                                                                                                                                          linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Diskette
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                                                                                                            9.1%; Score 12; DB 2;
100.0%; Pred. No. 4.4e+
tive 0; Mismatches
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METHOD OF ASSAYING DIFFERENTIAL
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                                                                                                                                        Length 26;
                                                                                                               Indels
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Sequence 3, Application US/08486857 Patent No. 6075181

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GENERAL INFORMATION:

Kucherlapati,

Raju

APPLICANT: APPLICANT:

APPLICANT:

Jakobovits, Aya Klapholz, Sue Brenner, Daniel G

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US-08-724-752-19/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; MOLECULE TYPE: DNA (genomic)
US-08-486-857-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Kucherlapati, Raju
APPLICANT: Jakobovits, Aya
APPLICANT: Brenner, Daniel G.
APPLICANT: Capon, Daniel J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEX: 66141 PENNIE INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING FACTOR 435
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Halluin, Albert P.
REGISTRATION NUMBER: 25,227
REGISTRATION NUMBER: 7639-042
                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DDS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/486,857
FILING DATE: 07-UN-1995
                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: 415-843-3660
                                                                                                                                                                                                                                                           APPLICANT: Klaphoz, Sue
TITLE OF INVENTION: HUMAN ANTIBODIES DERIVED FROM IMMUNIZED
TITLE OF INVENTION: XENOMICE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
PRRENT APPLICATION DATA:
                                                                                                                                                                                              STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY:
                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS:
                                                                                                                                                                                                            ADDRESSEE:
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                                                                                                                         10020
                                                                                                                                                                          T: 1251 Avenue of the Americas New York
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6150584
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Y: U.S.A.
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                                                                                                                                          USA
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                                                                                                                                                                                                            FISH & NEAVE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      9.1%; Score 12;
100.0%; Pred. No.
tive 0; Mismatcl
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US/08/724,752
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hes 0;
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RESULT 55
US-09-225-928-358/c
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GENERAL INFORMATION:
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Best Local (
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SEQUENCE CHARACTERISTICS:
                                                                TELEFAX: 415-854-0875
INFORMATION FOR SEQ ID NO: 358:
SEQUENCE CHARACTERISTICS:
                                                                                                                           COMPUTER: IBM COMPATIBLE
COPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/225,928
FILING DATE: 05-0an-1999
CLASSIFICATION CUNKNOWN>
PRIOR APPLICATION NUMBER: 08/859,998
APPLICATION NUMBER: 08/859,998
FILING DATE: 21-MAY-1997
ATTORNEY/ACENT INFORMATION:
NAME: Field, Bret E.
REGISTRATION NUMBER: 37,620
REFERENCE/DOCKET NUMBER: 09096/002001
TELECOMMUNICATION INFORMATION:
TTELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 29-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Haley Jr., James F
REGISTRATION NUMBER: 27,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: 1i
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 2200 Sand Hill Road, Suite
                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Chenchik, Alex
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: METHOD OF ASSAYING DIFFERENTIAL
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TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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ZIP: 94025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Menlo Park
                                                                                                                     TELEPHONE: 415-322-5070
                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE:
                                                 ENGTH: 26 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                               Diskette
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100.0%; Pred. No. 4.
ive 0; Mismatches
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SEQUENCE DESCRIPTION: SEQ ID NO: 358:
US-09-225-201B-358
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OTHER INFORMATION: oligonucleotide primer;
SEQUENCE DESCRIPTION: SEQ ID NO: 358;
US-09-225-928-358
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US-09-225-201B-358/c
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GENERAL INFORMATION:
                                                                            Query Match
Best Local Similarity
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version:
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                  NAME: Field, Bret E.
REGISTRATION NUMBER: 37,620
REFERENCE/DOCKET NUMBER: 09096/002001
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/859,998 FILING DATE: 21-MAY-1997 ATTORNEY/AGENT INFORMATION:
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ADDRESSEE: Fish & Richardson, P.C.
STREET: 2200 Sand Hill Road, Suite 100
CITY: Menlo Park
STATE: CA
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Bibilashvilli, Robert
TITLE OF INVENTION: METHOD OF ASSAYING DIFFERENTIAL
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                                                                                                                                                                           FEATURE:
                                                                                                                                                                                          MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
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                               109
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19 TGTCCTACCTGC 8
                                                               12;
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                     TGTCCTACCTGC 120
                                                                                                                                                                                                                    LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/09/225,201B FILING DATE: 05-Jan-1999 CLASSIFICATION: <Unknown>
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                                                                                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                      ELEPHONE: 415-322-5070
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100.0%; Pred. No. 4.
ive 0; Mismatches
                                                                          9.1%; Score 12;
100.0%; Pred. No.
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                                                               Mismatches
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                                                           4.4e+03;
hes 0;
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RESULT 58
US-09-614-092A-19/c
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                                                           PRIOR APPLICATION NUMBER: 08/724,752
PRIOR FILING DATE: 1996-10-02
PRIOR APPLICATION NUMBER: 08/430,938
PRIOR FILING DATE: 1995-04-27
PRIOR APPLICATION NUMBER: 08/234,145
PRIOR FILING DATE: 1994-04-28
PRIOR APPLICATION NUMBER: 08/12,848
PRIOR APPLICATION NUMBER: 08/212,848
PRIOR FILING DATE: 1993-08-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR FILING DATE: 1994-04-28
PRIOR APPLICATION NUMBER: 08/112,848
PRIOR FILING DATE: 1993-08-27
PRIOR APPLICATION NUMBER: 08/031,801
PRIOR FILING DATE: 1993-03-15
PRIOR FILING DATE: 1993-03-19
PRIOR FILING DATE: 1992-07-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19, Application US/09614092A Patent No. 6713610
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                           TITLE OF INVENTION: HUMAN ANTIBODIES DERIVED FROM IMMUNIZED XENOMICE FILE REFERENCE: Cell 4.17 DIV2
CURRENT APPLICATION NUMBER: US/09/614,092A
CURRENT FILING DATE: 2000-07-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 07/610,515
PRIOR FILING DATE: 1990-11-08
PRIOR APPLICATION NUMBER: 07/466,008
PRIOR FILING DATE: 1990-01-12
NUMBER OF SEQ ID NOS: 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: CELL 4.8 FWC CPA
CURRENT APPLICATION NUMBER: US/08/923,138
CURRENT FILING DATE: 1997-09-16
                                                                                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: KUCHERLAPATI,
APPLICANT: JAKABOVITS, J
APPLICANT: BRENNER, DAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 08/430,938
PRIOR FILING DATE: 1995-04-27
PRIOR APPLICATION NUMBER: 08/234,145
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 26
TYPE: DNA
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TITLE OF INVENTION: HUMAN ANTIBODIES DERIVED FROM IMMUNIZED XENOMICE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
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                   APPLICATION NUMBER: 08/031,801
FILING DATE: 1993-03-15
PPLICATION NUMBER: 07/919,297
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                                                                                                                                                                                                                                                                                                                                           CAPON, DANIEL J.
KLAPHOLZ, SUE
                                                                                                                                                                                                                                                                                                                                                                                        JAKABOVITS, AYA
BRENNER, DANIEL G.
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KLAPHOLZ, SUE
BRENNER, DANIEL G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        9.1%; Score 12;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4.4e+03;
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PRIOR FILING DATE: 1990-01-12

PRIOR APPLICATION NUMBER: 07/610,515

FILING DATE: 1990-11-08
APPLICATION NUMBER: 07/466,008

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RESULT 59
US-08-911-894-25/c
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                                                        US-08-911-894-25
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 Query Match
Best Local Similarity
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LENGTH: 26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Saxon, Andrew
APPLICANT: Zhang, Ke
APPLICANT: Elijieda, Shigeharu
APPLICANT: Fujieda, Shigeharu
TITLE OF INVENTION: IMMUNOGLOBULIN TRANS-SPLICED TRANSCRIPTS
TITLE OF INVENTION: AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: PCT/US96/05928
PRIOR FILING DATE: 1996-04-29
NUMBER OF SEQ ID NOS: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                       TELEFAX: (512) 499-629
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Description of Artificial Sequence: Primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: MayField, Denise L.
REGISTRATION NUMBER: 33,732
REFERENCE/DOCKET NUMBER: 43496.0006
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 19-AUG-1
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDATEST: OF STREET: OF AUStin
                                                                                      STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/911,894
FILING DATE: Concurrently Herewith
                                                                          TOPOLOGY:
                                                                                                                                                                                            TELEPHONE:
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                                                                                                                         ENGTH:
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12; Conserv
                                                                                                     nucleic acid
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816 Congress Avenue, Suite 1900
                                                                                                                         27 base pairs
                                                                                                                                                                            : (512) 499-6200
(512) 499-6290
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                        linear
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                                                                                      single
   9.1%;
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Pred. No. 4.4e+03;
Score 12; DB 3; I
Pred. No. 4.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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                  Length
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                   27;
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APPLICANT: MUKUMOTO, Fujio
APPLICANT: NISHIO, Shoichi
APPLICANT: NISHIO, Shoichi
APPLICANT: NISHIO, Shoichi
APPLICANT: MITSUDA, Satoshi
TITLE OF INVENTION: DNA Fragments Containing Biotin Biosynthetase Gene and
TITLE OF INVENTION: Use of the Same
FILE REFERENCE: 0152-0490P
CURRENT APPLICATION NUMBER: US/09/180,109A
CURRENT FILING DATE: 1998-12-03
PRIOR APPLICATION NUMBER: 09/047838 JAPAN
PRIOR APPLICATION NUMBER: 09/047838 JAPAN
PRIOR FILING DATE: 1997-03-03
NUMBER OF SEQ ID MOS: 52
SOOTWARE: Patentin Ver. 2.0
LENGTH: 28
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US-09-180-109A-47/c
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                                                                             ; OTHER INFORMATION: NETEX9 56 US-09-304-232-669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-304-232-669/c
                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/304,232
CURRENT FILING DATE: 1999-05-03
EARLIER APPLICATION UNMBER: US 60/084,641
EARLIER FILING DATE: 1998-05-07
NUMBER OF SEQ ID NOS: 909
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 669
LENGTH: 29
LENGTH: 29
                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Fan, Jian Bing
APPLICANT: Chakravarti, Aravinda
APPLICANT: Halushka, Marc Kenneth
APPLICANT: Case Western Reserve University School of Medicine
APPLICANT: Affymetrix, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 669, Application US/09304232 Patent No. 6525185
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                                           Query Match
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                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Polymorphisms Associated With TITLE OF INVENTION: Hypertension FILE REFERENCE: 018547-034210US
                                                                                                                       TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence FEATURE:
                                                                                                                   FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
                    Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12;
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       12;
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                      Similarity
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       Conservative
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100.0%; Pr
                    9.1%; Score 12; DB 4;
100.0%; Pred. No. 4.4e+0
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Pred. No.
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                      4.4e+03;
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                                         Length 29;
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RESULT 63
US-07-714-131C-27
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US-07-931-473B-27
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                                                                                                                                                                                                                                                  Sequence 27, Application US/07714131C Patent No. 5475096
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 27, Application US/07931473B
                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPAX: (303) 850-9401
INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
LENGTH: 30 nucleotides
TYPE: NUCLEIC ACID
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 5.25 inch, 360 Kb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM compatible
OPERATING SYSTEM: MS-DOS
SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/931,473B
                                                                                                                                       APPLICANT: Larry Gold
APPLICANT: Craig Tuerk
TITLE OF INVENTION: Nucleic Acid Ligands
NUMBER OF SEQUENCES: 344
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (303) 850-9900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
ATTORNEY/AGENT INFORMATION:
NAME: Barry J. Swanson
REGISTRATION NUMBER: 33,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 5
COMPUTER: IBM compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Larry Gold
APPLICANT: Craig Tuerk
TITLE OF INVENTION: Nucleic Acid Ligands
NUMBER OF SEQUENCES: 335
CORRESPONDENCE ADDRESS:
ADDRESSEE: Beaton & Swanson, P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS: sir
TOPOLOGY: linear
                                                                                                          STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 19
CLASSIFICATION:
                                       ZIP: 80237
                                                                               STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 80237
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Denver
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET:
                                                        COUNTRY:
                                                                                                                              ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                               Local
                                                                                                                                                                                                                                                                                                                                                                             91 ATGGAGCGCTCG 102
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                                                                                                                                                                                                                                                                                                                                                                                                                           ch 9.1%; Score 12; DB 1; Similarity 83.3%; Pred. No. 4.4e+03; 10; Conservative 2; Mismatches 0
                                                                                            Denver
                                                                                                                                                                                                                                                                                                                                                          5 AUGGAGCGCUCG 16
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                                                                            Colorado
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                                                                                                            4582 South Ulster Street Parkway, #403
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4582 South Ulster Street Parkway, #403
                                                           USA
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                                                                                                                           Beaton & Swanson, P.C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     360 Kb storage
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US-08-412-110-27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 27, App. Patent No. 56706
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Best Local:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                   APPLICATION UNMBER: US/08/412
APPLICATION UNMBER: US/08/412
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 07/714.13
APPLICATION UNMBER: 07/536,42
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 07/536,42
PILING DATE: June 11, 1990
ATTORNEY/AGENT INFORMATION:
NAME: BEATY J. SWANBON
REGISTRATION NUMBER: 33,215
REFERENCE/DOCKET NUMBER: 33,215
REFERENCE/DOCKET NUMBER: NEXO
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                      TELEFAX: (303) 793-3433 INFORMATION FOR SEQ ID NO: 27:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (303) 850-9900
TELEPAX: (303) 850-9401
INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/07/71
FILING DATE: June 10, 1991
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
ATTORNEY/AGENT INFORMATION:
NAME: Barry J. Swanson
REGISTRATION NUMBER: 33,215
                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Larry Gold APPLICANT: Craig Tuerk TITLE OF INVENTION: Nu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM: MS-DOS
SOFTWARE: $ WordPerfect 5.1
CURRENT APPLICATION DATA:
                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER:
TELEPHONE: (303) 850-9900
                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM: MS-DOS
SOFTWARE: WordPerfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER:
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Colorado
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Application US/08412110
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                                                                                                                                                                                                                                                                                                                                                                                       IBM compatible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Swanson & Bratschun, L.L.C.
                                                                                                                                                                                                                                                                                                                                                                                                       Diskette, 3.50 inch, 1.44 MG storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             9.1%;
83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic Acid Ligands
                                                                                                                                                                                                              07/536,428
                                                                                                                                                                                                                                                                 07/714,131
                                                                                                                                                                                                                                                                                                                  US/08/412,110
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                                                                                                                         NEX01/C2
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Pred. No. 4.4e+03;
2; Mismatches (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Suite 200
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RESULT 66
US-08-469-609A-27
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US-08-409-442A-27
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 Sequence 27, Applicat
Patent No. 5843653
GENERAL INFORMATION:
APPLICANT: Larry
                                                                                                                                                                                      Matches
                                                                                                                                                                                                  Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 9.1%;
Best Local Similarity 83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 27, Application US/08409442A
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                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/714,131
FILING DATE: June 10, 1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/536,428
FILING DATE: June 11, 1990
ATTORNEY/AGENT INFORMATION:
NAME: BALTY J. SWANBOR
REGISTRATION NUMBER: 33,215
REFERENCE/DOCKET NUMBER: NEXO1/C3
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent No. 5696249
                                                                                                                                                                                                                                                                                                        TELEFAX: (303) 793-3433
INFORMATION FOR SEQ ID NO: :
SEQUENCE CHARACTERISTICS:
LENGTH: 30 nucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: Bauu _
CITY: Englewood
CTATE: Colorado
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Larry Gold
APPLICANT: Craig Tuerk
TITLE OF INVENTION: Nucleic Acid Ligands
NUMBER OF SEQUENCES: 374
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM compatible OPERATING SYSTEM: MS-DOS SOFTWARE: WordPerfect 5.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: C
                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE:
                                                                                                                                         91 ATGGAGCGCTCG 102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             91 ATGGAGCGCTCG 102
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                                              Application US/08469609A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         E: Swanson & Bratschun, L.L.C.
8400 E. Prentice Avenue, Suite
                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                             linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Diskette,
                                                                                                                                                                                                  9.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US/08/409,442A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3.50 inch,
                                                                                                                                                                                   Score 12; DB 1;
Pred. No. 4.4e+03;
2; Mismatches (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Suite 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.44 MG storage
                                                                                                                                                                                                                Length 30;
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US-09-143-190-27
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US-08-469-609A-27
                                                                                                                                                                                                                                                                                                                                       Matches
              COUNTRY:
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                                                           STREET:
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Sequence 27, Application US/09143190 Patent No. 6110900
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Best Local Similarity 83.3%;
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FILING DATE: June 6, 1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/428,964
FILING DATE: April 25, 1995
PRIOR APPLICATION NUMBER: 08/412,110
APPLICATION MATCH 27, 1995
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION UMBER: 08/409,442
APPLICATION DATA:
APPLICATION MATCH 24, 1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                         GENERAL INFORMATION:
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TELEFAX: (303) 793-343:
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERITICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:

NAME: BATTY J. SWAIISON
REGISTRATION NUMBER: 33,215
REFERENCE/DOCKET NUMBER: NEXO
TELECOMMUNICATION INFORMATION:
TELEPHONE: (303) 793-3333
                                                        ADDRESSEE: Swanson &
                                                                                                    APPLICANT: Larry Gold
APPLICANT: Craig Tuerk
TITLE OF INVENTION: Nucleic
NUMBER OF SEQUENCES: 374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 07/714,131 FILING DATE: June 10, 1991 PRIOR APPLICATION DATA:
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TITLE OF INVENTION: Nucleic Acid Ligands
NUMBER OF SEQUENCES: 374
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
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                 CITY: Englewood
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nucleic acid
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Colorado
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                                       B: Swanson & Bratschun, L.L.C.
8400 E. Prentice Avenue, Suite
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8400 E. Prentice Avenue, Suite
                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 12; DB 2; Pred. No. 4.4e+03;
                                                                                                                          Acid Ligands
                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                         Suite 200
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US-09-502-344-27
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                                                                                                                                                                                                                                                                                                                        Patent No. 6331398
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 08/409,44:
FILING DATE: March 24, 1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/714,13:
FILING DATE: June 10, 1991
PRIOR APPLICATION NUMBER: 07/536,42:
APPLICATION NUMBER: 07/536,42:
FILING DATE: June 11, 1990
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 08/469,609
FILING DATE: June 6, 1995
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 08/428,964
FILING DATE: April 25, 1995
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 08/412,110
FILING DATE: March 27, 1995
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (303) 793-343
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                               ZIP: BULL.
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.44 MG storage
COMPUTER: IBM compatible
COMPUTER: MS DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER: 33,215
REFERENCE/DOCKET NUMBER: NEXO1/C6
TELECOMMUNICATION INFORMATION:
TELEPHONE: (303) 793-3333
                                                                                                                                              STREET: 8400
CITY: Englewood
STATE: Colorado
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PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                             APPLICANT: Larry Gold
APPLICANT: Craig Tuerk
TITLE OF INVENTION: Nucleic Acid Ligands
NUMBER OF SEQUENCES: 374
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3
COMPUTER: IBM compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Match 9.1%;
Local Similarity 83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH:
APPLICATION NUMBER: US/09/502,344
                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                  ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                            27, Application US/09502344
o. 6331398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          91 ATGGAGCGCTCG 102
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                  ARE: WordPerfect 8.0 APPLICATION DATA:
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                                                                                                                                                                                              E: Swanson & Bratschun, L.L.C.
8400 E. Prentice Avenue, Suite
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 12; DB 3; 1
Pred. No. 4.4e+03;
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                                                                                                                                                                                                Suite 200
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 30;
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OTHER INFORMATION: Description of Artificial Sequence: Primer US-09-937-832-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-502-344-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-937-832-16/c
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Best Local S
Matches 10
                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 16, Application US/09937832
Patent No. 6586200
GENERAL INFORMATION:
Query Match
Best Local Similarity
                                                                                                                                                                        SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 16
                                                                                                                                                                                                       APPLICANT: OMURA, SATOSHI
APPLICANT: ABE, AXIO
TITLE OF INVENTION: METHOD FOR DETECTING SUBSTANCES INHIBITING THE
TITLE OF INVENTION: BACTERIAL TYPE III SECRETION MECHANISM AND FUNCTION OF
TITLE OF INVENTION: SECRETORY PROTEINS THEREOF
FILE REFERENCE: KP-9106
CURRENT APPLICATION NUMBER: US/09/937,832
CURRENT FILING DATE: 2002-05-21
PRIOR APPLICATION NUMBER: PCT/JP01/00377
PRIOR PILING DATE: 2001-01-22
NUMBER OF SEQ ID NOS: 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 30 nucleotiden
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: BAILY J. SWANSON
REGISTRATION NUMBER: 33,215
REFERENCE/DOCKET NUMBER: NEXC
TELECOMMUNICATION INFORMATION:
TELEPHONE: (303) 793-3333
                                                                                          TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/143,190
APPLICATION NUMBER: 27, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: June 10,
PRIOR APPLICATION DATA:
APPLICATION NUMBER: (
FILING DATE: June 11)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 08/428, FILING DATE: April 25, 1995 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: June 6, PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 08/409, APPLICATION NUMBER: 08/409, FTI.ING DATE: March 24, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: 08/412, PILING DATE: March 27, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 07/714 FILING DATE: June 10, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               91 ATGGAGCGCTCG 102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5 AUGGAGCGCUCG 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   June 11, 1990
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 9.1%;
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 Score 12; DB 4; Length 30; Pred. No. 4.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 12; DB 3;
Pred. No. 4.4e+03;
2; Mismatches (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 3; Length 30;
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APPLICANT: Abbott Laboratories
APPLICANT: Scheffel, C.
APPLICANT: Scheffel, C.
APPLICANT: Scheffel, C.
TITLE OF INVENTION: Nucleic Acid Primers And Probes For
TITLE OF INVENTION: Detecting Tumor Cells
FILE REFERENCE: 6589 US.01
CURRENT APPLICATION NUMBER: US/09/419,212
CURRENT FILING DATE: 1999-10-15
NUMBER OF SEQ ID NOS: 8
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 4, Application US/09419212

Patent No. 6203992

GENERAL INFORMATION:

APPLICANT: Abbott Laboratories

APPLICANT: Scheffel, C.

APPLICANT: Granados, E.

TITLE OF INVENTION: Nucleic Acid Primers And Probes For

TITLE OF INVENTION: Detecting Tumor Cells

FILE REFERENCE: 6589.US.01

CURRENT APPLICATION NUMBER: US/09/419,212

CURRENT APPLICATION NUMBER: US/09/419,212

CURRENT APPLICATION STORE SERVING SERVI
              RESULT 72
US-09-060-299-454/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-419-212-6/c
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US-09-419-212-4
                                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: Probe US-09-419-212-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: Probe US-09-419-212-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 6, Application US/09419212
Patent No. 6203992
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 15
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                          11 CCAGGACCTGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15 CCAGGACCTGC 25
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                                                                                                                                                                               15 CCAGGACCTGC 25
                                                                                                                                                                                                                                     11;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8.3%; Solitarity 100.0%; If Conservative 0;
                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                            8.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 11; DB 3; Le
; Pred. No. 1.5e+04;
0; Mismatches 0;
                                                                                                                                                                                                                                                            Score 11; DB 3;
; Pred. No. 1.5e+(
                                                                                                                                                                                                                                   <u>,</u>
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                                                                                                                                                                                                                                     Mismatches
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                                                                                                                                                                                                                                                                                                                                            RESULT 73
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APPLICANT: Hey, Patricia
APPLICANT: Kawaguchi, Yoshihiko
APPLICANT: Kawaguchi, Yoshihiko
APPLICANT: Metzker, Michael L
TITLE OF INVENTION: No. 6545137el Receptor
NUMBER OF SEQUENCES: 455
CORRESPONDENCE ADDRESS:
ADDRESSER: Niver-
                                                                                                                                                                                                                          Sequence 454, Application US/09402923A Patent No. 6555654
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Loc
Matches
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 60/043,553
FILING DATE: 15-APR-1997
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 60/048,740
FILING DATE: 05-JUN-1997
ATTORNEY/AGENT INFORMATION:
NAME: B.J.Sadoff
REGISTRATION NUMBER: 36,663
REFERENCE/DOCKET NUMBER: 620-35
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (703)816-409
TELEFAX: (703)816-4100
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/060,299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 16 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 15
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 1100 No. CITY: Arlington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                    APPLICANT: Todd, John A
Hess, John W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           109 TGTCCTACCTG 119
                                                                                                                                                                                                                                                                                                                                                                                                                           14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        E: Nixon and Vanderhye
1100 No. 6545137th Glebe Road, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Caskey, Charles
Cox, Roger D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Todd, John A
Hess, John W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
Caskey, Charles T
Cox, Roger D
Gerhold, David
Hammond, Holly
Hey, Patricia
Kawaguchi, Yoshihiko
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (703)816-4091
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              8.3%; Score 11; DB 4; L
100.0%; Pred. No. 1.5e+04;
htive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Release #1.0, Version #1.25 (EPO)
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US-07-990-297-13
                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: GROSZ, RON
APPLICANT: JENSEN, WARK A
TITLE OF INVENTION: IMPROVEI
TITLE OF INVENTION: AMPLIFIC
TITLE OF INVENTION: SEGMENT;
TITLE OF INVENTION: MESTED
NUMBER OF SEQUENCES: 22
                                                                                                                                                                                                                                                                                                 Patent No. 53*v.__
Patent No. 53*v.__
PATENT INFORMATION:
PROSZ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (703)816-4100
INFORMATION FOR SEQ ID NO: 454:
SEQUENCE CHARACTERISTICS:
COMPUTER READABLE FORM: MEDIUM TYPE: Diskette,
                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: E. I. du Pont de Nemours and Company STREET: 1007 Market Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 100 les 11; Conservative
                                                                                                 CITY: Wilmington
                                                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: 36,663
REFERENCE/DOCKET NUMBER: 62
TELECOMMUNICATION INFORMATION:
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TITLE OF INVENTION: No. 6555654el LDL-Receptor
NUMBER OF SEQUENCES: 455
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE DESCRIPTION: SEQ ID NO: 454:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             109 TGTCCTACCTG 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                          14 TGTCCTACCTG 4
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PILING DATE: 14-Feb-2001
IOR APPLICATION DATA:
APPLICATION NUMBER: PCT/GB98/01102
FILING DATE: 15-APR-1998
APPLICATION NUMBER: US 60/043,553
FILING DATE: 15-APR-1997
APPLICATION NUMBER: US 60/048,740
FILING DATE: 05-JUN-1997
IORNEY,AGENT INFORMATION:
                                            19898
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Arlington
STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Nixon and Vanderhye STREET: 1100 No. 6555654th Glebe Road, Eighth Floor
                                                                               Delaware
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (703)816-4091
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: B.J.Sadoff
                                                                                                                                                                                                                                                                                                                                                        Application US/07990297
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                                                             U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; F1
                                                                                                                                                                         IMPROVED METHOD FOR
AMPLIFICATION OF TARGETED
SEGMENTS OF NUCLEIC ACID USING
NESTED POLYMERASE CHAIN REACTION
22
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100.0%; Pred. No. 1.9
1ve 0; Mismatches
  3.50 inch, 1.0 MB
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US-08-373-124A-1749/c
; Sequence 1749, Application US/08373124A
; Patent No. 5646042
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No. GENERAL II
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Best Local Similarity 100.0%;
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION UNMER: US/08/373,124A
FILING DATE: January 13, 195
PRIOR APPLICATION UNMER: 08/245,466
FILING DATE: May 18, 1994
APPLICATION NUMBER: 08/192,943
FILING DATE: February 7, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: February 7, 1992
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/986,422
APPLICATION HOMBER: 07/987,132
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INFORMATION FOR SEQ ID NO:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/990
FILING DATE: 19921209
CLASSIFICATION: 435
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
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LENGTH: 17 base pairs
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NAME: GEIGER, KATHLEEN W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM CONTROL OPERATING SYSTEM:
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STREET: Suite 4700
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McSwiggen, James
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CANCER USING RIBOZYMES
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100.0%; Pred. No.
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REFERENCE/DOCKET NUMBER: 209/035
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (213) 489-1600
; TELEPAX: (213) 955-0440
; TELEPAX: (213) 955-0440
; TELEPAX: (213) 955-0440
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO: 1749:
SEQUENCE CLARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
stranupeDNESS: single
TOPOLOGY: linear
US-08-373-124A-1749

Query Match
Best Local Similarity 100.0%; Pred. No. 1.5e+04;
Matches 11; Conservative 0; Mismatches 0; Indels 0; Gaps
Qy 91 ATGGAGCGCTC 101
Db 12 ATGGAGCGCTC 101
Search completed: February 2, 2005, 23:37:50
Job time: 14.6615 secs
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C 1 13 2.6 21 8 AZ873829 C 2 13 2.6 26 8 AZ762572 C 4 13 2.6 28 1 AU25572 C 5 13 2.6 28 1 AU25572 C 6 6 12 2.4 22 8 AZ364675 C 7 12 2.4 22 8 AZ304010 C 7 12 2.4 23 8 AZ36557 C 10 12 2.4 26 8 AZ452570 C 10 12 2.4 26 8 AZ452570 C 11 12 2.4 28 1 AA906876 C 11 12 2.4 29 8 AZ421270 C 13 12 2.4 29 8 AZ421270 C 13 12 2.4 29 8 AZ623010 C 14 11 2.2 12 6 CF305550 C 16 11 2.2 12 6 CF30693 C 17 11 2.2 12 16 AI017839 C 18 11 2.2 19 8 AZ641152 C 18 11 2.2 19 8 AZ641152 C 19 11 2.2 19 8 AZ648404 C 21 11 2.2 22 1 AA9989077 C 22 11 2.2 22 1 AA9890077 C 22 11 2.2 24 8 AZ331171 C 24 11 2.2 24 AZ6 AZ531171	Post-processing: Listing first 150 summaries EST:* 1: gb_est1:* 2: gb_est2:* 3: gb_htc:* 4: gb_est3:* 5: gb_est4:* 6: gb_est5:* 6: gb_est5:* 7: gb_est6:* 8: gb_gss1:* 9: gb_gss2:* Pred. No. is the number of results predicted by classore greater than or equal to the score of the reand is derived by analysis of the total score dispensed by analysis of the SumMARIES Result Query No. Score Match Length DB ID	Word size: 0 Total number of hits satisfying chosen parameters: Minimum DB seg length: 0 Maximum DB seg length: 30	Title: US-10-048-046-1_COPY_1516_2013 Perfect score: 498 Sequence: 1 tgccctctgcaaggaagccagctactg. Scoring table: OLIGO_NUC Gapop 60.0 , Gapext 60.0 Searched: 32822875 seqs, 18219865908 residues	GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compug etc - nucleic search, using sw model February 2, 2005, 15:07:30; Searc
AZ873829 ZMO187C15 BZ356062 SALK 1280 AZ762572 1M0557P02 AU255721 AU255721 AU255721 AU255721 AZ364675 1M00110K20 AZ304010 1M0003023 AZ795667 ZM0029J15 CCF303555 ABF102- AZ436027 1M0233C06 AA906876 c120e05.8 AZ41270 1M0199022 AZ595520 1M0408M09 AZ633010 1M0460E14 CCF306933 HDA105- AI017839 0V008F12.x BM400383 5009-0-72 AZ361152 1M0104A16 AZ643124 1M0506P02 AZ648104 1M0517123 AZ688404 1M0517123 AZ995149 ZM0280D22 AA989077 or88e03.8 AL490359 T. brucei CD531171 09120 Aza AZ331594 1M0059D12	o have a eing printed, on.	0 000	gctactggggcgtaactgc 498 cc	en Ltd. c c h time 1529.28 Seconds alignments) c alignments c 1 Million cell updates/sec
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1625 AU013625 1662 AU013625 1714 AU013714 1872 E012766-0 1872 E01276600 1872 E01276600 1872 E01276600 1872 E01276600 1872 E01276600 1872 E01276000 1872 E0127600000000000000000000000000000000000	- 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	5 5009-0-4 5 E012845- 1 ub33a03. 6 DKFZp434 5 5009-0-1 3 13104 Ar 9 CLD101	.លលល់ទំល់សិ <i>កី</i> ប៉	\$ qa01e11. \frac{1}{2} \text{qa01e11.r1} \text{y124e12.r1} \text{yr08g12.r1} \text{01119027E} \text{3119027E}

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AZ873829
21 l
2M0187C15R Mouse 10kb plasmid
Clone UUGC2M0187C15 R, genomic
        Dunn, D., Aoyagi, A.,
Islam, H., Longacre, S
                              Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia; 1 (bases 1 to 21)
                                                                Mus musculus
                                                                                        AZ873829.1
GSS.
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                                                                                                                                                                                                                                           Longacre, S.,
                                                                                                    GI:13082289
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                                                                             mouse)
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AZ315933
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AZ4708673
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AJ646986
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AZ582153
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       Mahmoud, M.,
                                                                                                                          genomic
                                                                                                                                                                                                          ALIGNMENTS
                                                                                                                           d UUGCIM library Mus musculus genomic
                                        Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                         survey sequence.
                  Beacorn, T.,
    Meenen, E., Pedersen, T.,
                  Duval, B., Hamil, C.,
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AJ797003 AJ797003
BQ592774 E012124-0
CA587523 LBE13p44P
R89803 yP91b12.r1
AZ346865 1M0082J15
AZ419335 1M0155J17
AZ759839 1M0553B02
AZ828697 ZM0105015
AZ944762 ZM0205N19
BH900958 KG07822-3
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IMO553P10
IMO558P17
D IMO569K11
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7 2M0085E05
2 ot53d06.s
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5 1M0514N12
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1M0250A16
1M0315H02
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AJ791120
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                                                                                      ACCESSION
VERSION
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       REFERENCE
                                                                         KEYWORDS
                                                                                                                                  DEFINITION
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Best Local S
Matches 13
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                                                                                                                                                                                                                    132 CCTGTACTGGGGC 144
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Arabidopsis thaliana (thale cress)
Arabidopsis thaliana
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
1 (bases 1 to 24)
                                                                                                                   BZ356062 24 bp DNA linear GSS SALK 128022.33.70.x Arabidopsis thaliana TDNA insertion Arabidopsis thaliana genomic clone SALK 128022.33.70.x,
                                                                                               survey sequence.
BZ356062
                                                                                      BZ356062.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0187 row: C column: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R. Mouse whole genome scaffolding with paired end reads from plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       University of Utah (University of Utah Rm. 308, Biomedical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (2000)
Contact: Robert B.
                                                                                                                                                                                                                                                              Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                  ilarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  quality sequence stop: 21.
                                                                                                                                                                                                                                                                                                     was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pMD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /lab_host="E. Coli strain XL10-Gold, T1-resistant, /Clone_lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: pW042nv; Purified genomic DNA from the Jac
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mol_type="genomic DNA
/strain="C57BL/6J"
/db_xref="taxon:10090"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sex="Male"
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                                                                                    GI:24947395
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Pred. No.
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Mismatches 0;
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The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
  Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Brror: (
Plate: 0557 row: P column: 02
Seg primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                          University of Utah Genome Center University of Utah Rm. 308, Biomedical Polymers Res
                                                                                                                                                                                                                            Unpublished (2000)
Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                         Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Wahmoud, M., Weenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. Mouse whole genome scaffolding with paired end reads from 10kb Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 26)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AZ762572

26 bp DNA linear GSS 16-FEB-2001
1M0557P02R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0557P02 R, genomic survey sequence.
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Unpublished (2001)
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A Sequence-Indexed Library of Insertion Mutations in the
                                                                                           Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                         plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AZ762572.1 GI:12872712
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Joseph R. Ecker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mus musculus (house mouse)
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llarity 100.0%;
Conservative (
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                                                                                                                                            USA
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/db_xref="taxon:3702"
/clone="SALK_128022.33.70.x"
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|mol_type="genomic DNA"
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                                            Std Error: 0.00
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                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata Mammalia; Eutheria; Rodentia; Sciurogr 1 (bases 1 to 28)
Xato, K. and Matoba, R.
Generation of expressed sequence tags Unpublished (2002)
                                                                                                                                                                                                Contact: Kikuya Kato
Graduate School of Biological Sciences
Nara Institute of Science and Technology
8916-5 Takayama, Ikoma, Nara 630-0101, Ja
Tel: 81-743-72-5589
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Location/Qualifiers
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AU255721 3'-directed mouse cDNA library
BED0006262 3', mRNA sequence.
                                                                                                                              Email: kkato@bs.aist-nara.ac.jp,
URL:http://love2.aist-nara.ac.jp/BED/index.html.
Location/Qualifiers
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                     /mol_type="mRNA"
/db_xref="taxon:10090"
/clone="BED0006262"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Laboratory Mouse DNA Resource
/tissue_type="brain"
                                                                                         organism="Mus musculus"
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strain="C57BL/6J"
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R. Mouse whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert B. Weiss University of Utah Genome Center University of Utah
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larity 100.0%; Pred. No. 1.8e+05;
Conservative 0; Mismatches 0;
                                                     (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA pollymerase and T4 pollymcleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"/clone_lib="Mouse_10kb_plasmid_UUGCIM_library"/note="Vector: PWD42nv; Purified_genomic_DNA_from_M.musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtai
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strain="C57BL/6J"
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Location/Qualifiers
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University of Utah Genome Center
University of Utah
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Insert Length: 10000 Std Error: (
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Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AZ304010 22 bp DNA linear GSS 29-SEP-200
1M0003023R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AZ304010.1 GI:10339557
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ass: plasmid ends
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        308,
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD42 (gil4732114 |gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coli XL10-Gold (Stratagene) cells
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/Clone_lib="Mouse 10kb plasmid UUGC1M library"/note="Vector: PW042nv; Purified genomic DNA from M.musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   clone="UUGC1M0003023"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%;
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AZ785667
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VERSION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts plasmid inserts Contact; Robert B. Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0029 row: J column: 15
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 23)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AZ785667
AZ785667.1 GI:12922656
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            University of Utah Genome Center University of Utah
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Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polymucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD42 (gi|4732114|gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coli XL10-Gold (Stratagene) cells
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/Clone_lib="Mouse 10kb plasmid UUGCIM library"/note="Vector: pW042nv, Purified genomic DNA from M.musculus C57BL/6J (male) was obtained from the Jackson
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sex="Male"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2.4%;
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Pred. No.
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6.3e+05;
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                                                                                                              RESULT 9
AZ436027/c
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                                                                                   DEFINITION
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                                           AZ436027 26 bp DNA linear GSS 03 1M0233C06R Mouse 10kb plasmid UUGC1M library Mus musculus clone UUGC1M0233C06 R, genomic survey sequence.
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Le Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H. Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABF1--02-J13.g1 ABF3-overexpressing transgenic rice lambda cDNA library (ABF1) Oryza sativa (japonica cultivar-group) clone ABF1--02-J13, mRNA sequence.
                    AZ436027
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta;
Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         œ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Nahm B.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ehrhartoideae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /clone_lib="ABF3-overexpressing transgenic rice lambda phage cDNA library (ABF1)"
/note="Vector: pBluescript SK(+); Site_1: EcoRI; Site_2: XhoI; Leaf was dried for 2hrs. cDNA was inserted into lamda Uni-2Ap Xr vector at 5' end with EcoRI and 3' end with XhoI site. mRNA was prepared from ABA-responsive element binding transcription factor 3 overexpression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        and selected for ampicillin resistance."
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/dev_stage="14 days after germination"
/lab_host="E_coli SOLR"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Oryza sativa (japonica cultivar-group)"
/mol_type="mRNA"
/cultivar="Nackdong"
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100.0%; Pred. No.
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100.0%; Pred. No.
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3. 6.4e+05;
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ORGANISM
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Best Local S
Matches 12
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AA906876 28 bp mRNA linear EST 10-JUN olloe05.s1 Soares NFL T GBC S1 Homo sapiens CDNA Clone IMAGE:1524032 3' similar to SW:Y032 HUMAN Q15034 HYPOTHETICAL PROTEIN KIAA0032. ;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0233 row: C column: 06
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Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            University of Utah Genome Center University of Utah
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Contact: Robert B.
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                                                                                                                                                                                                                  GTGGCTCTCCAG 324
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                                                                                                                                                                                                                                                                                                                                                      Laboratory Nouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWHPA2 (gi[4732114]gb]AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coll XL10-Gold (Stratagene) cells
and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /clone_lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: pWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           organism="Mus musculus"
/mol_type="genomic DNA"
/strain="C57BL/6J"
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'clone="UUGC1M0233C06"
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                                                                 EST 10-JUN-1998
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423 TCGGCAGAACAT 434
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                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 29)

Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29 bp
1M0199022F Mouse 10kb plasmid UU
clone UUGC1M0199022 F, genomic s
AZ421270
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This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (infc@image.llnl.gov) for further information.
Trace considered overall poor quality
Insert Length: 458 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham
High quality sequence stop: 1.
Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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1 (bases 1 to 28)
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                                                                                          plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AZ421270.1 GI:10545283
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone lib="Soares NFL T_GBC_S1"
/clone lib="Soares NFL T_GBC_S1"
/note="Organ: pooled; Vector: pT7T3D-Pac (Pharmacia) with
a modified polylinker; Site 1: Not I; Site 2: Eco RI;
Equal amounts of plasmid DNA from three normalized
libraries (fetal lung NbHL19W, testis NHT, and B-cell
NCI_CGAP_GCB1) were mixed, and ss circles were made in
vitro. Following HAP purification, this DNA was used as
tracer in a subtractive hybridization reaction. The driver
was PCR-amplified cDNAs from pools of 5,000 clones made
from the same 3 libraries. The pools consisted of
I.M.A.G.E. clones 297480-302087, 682632-687239,
726408-728711, and 729096-731399. Subtraction by Bento
Soares and M. Fatima Bonaldo. "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone="IMAGE:1524032"
/lab_host="DH108"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          genomic survey sequence.
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. 6.4e+05;
thes 0;
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AZ595520
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Best Local (
                                                                                                     TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   283 ACATGGAAAAAC 294
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18
                                               plasmid inserts
Unpublished (2000)
                                                                                                                       Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weise, R.
                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
(Dases 1 to 29)
                                                                                                                                                                                                                                                                                                       Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                            AZ595520 29 bp DNA linear GSS 13-DEC-200 IM0408M09F Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGCIM0408M09 F, genomic survey sequence.
AZ595520
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Insert Length: 10000 Std Error:
Plate: 0199 row: O column: 22
Seq primer: CGTTGTAAAACGACGGCCAGT
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Fax: 801 585 7177
Contact: Robert B. Weiss
University of Utah Genome Center
                                                                                                                                                                                                                                                                                                                                                                                    AZ595520.1 GI:11717710
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Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                               Mouse whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pDM42 (gi |4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid Rl. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/clone lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PWD42nv, Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mol_type="genomic DNA"
/strain="C57BL/6J"
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100.0%; Pred. No. 6.4
ive 0; Mismatches
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o. 6.4e+05;
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AZ623010/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        164 ACCCCTCCTCC 175
                                                                                           Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12;
                      plasmid inserts
Unpublished (2000)
Contact: Robert B.
                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1 (bases 1 to 30)
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                                                                                                                                                                                                                                                                                               Mus musculus
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Class: plasmid ends
High quality sequence stop: 29.
Location/Qualifiers
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Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0408 row: M column: 09
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Rm. 308, Biomedical
84112, USA
University of Utah
                                                                                                                                                                                                                                                                                                                    Mus musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymcleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel elettrophoresis. Vector DNA was prepared from a derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               clone="UUGC1M0408M09"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.4%;
  Genome Center
                         Weiss
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                                                                                                                                                                                                                                                                                                                                                                                                                                      30 bp DNA linear GSS 13-DEC-200
smid UUGC1M library Mus musculus genomic
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CF306933/c
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                                                                                                                                                                                                           Oryza sativa (japonica cultivar-group)
oryza sativa (japonica cultivar-group)
oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Spermatodeae; Oryzeae; Oryza.
     Genomics and Genetics Institute, of Bioscience and Bioinformatics,
                                                                                     1 (bases 1 to 12)

Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H.,

Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                             HDA1--05-E05.g1 OsHDAC1-overexpressing transgenic rice lambda phage cDNA library I (HDA1) Oryza sativa (japonica cultivar-group) cDNA clone HDA1--05-E05, mRNA sequence.
                                                             Contact: Nahm B.H.
                                                                                                                                                                                                                                                                                                                                                                       CF306933.1 GI:33678694
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Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0460 row: E column: 14
Seg primer: CGTTGTAAAACGACGGCCAGT
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Rm. 308, Biomedical Polymers Research Bldg.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pW142 (gi 4732114 |gb|AF129072.1), a copy-number inducible derivative of plasmid Rl. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          plasmid ends
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"/clone lib="Mouse 10kb plasmid UUGCIM library"/note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackso
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Laboratory Mouse DNA Resource
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 12; DB 8; Len
s; Pred. No. 6.5e+05;
GreenGene Biotech II
, MyongJi University
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                                Inc.; Division
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AI017839
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D.,
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12 AGAGCCTCGTG 2
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                                                                                                                                                                                      High quality sequence stop: 1.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                      cDNA Library Preparation: M. Bento Soares, Ph.D. cDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html
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16 bp mRNA linear EST 16-JUN-
0v08e12.x1 NCI_CGAP_Kid3 Homo sapiens cDNA clone IMAGE:1636750 ;
similar to TR:\(\bar{Q}\)35993 CYTOCHROME C OXIDASE III. ;contains
element L1 L1 repetitive element ;, mRNA sequence.
                                                                                                                                                                                                                                        Trace considered overall poor quality Seq primer: -40m13 fwd. ET from Amersham
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Tel: 82
Fax: 82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tumor Gene Index
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National Cancer Institute, Cancer Genome Anat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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AI017839.1 GI:3232175
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2 31 330 6193
2 31 321 6355
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Location/Qualifiers
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/lab host="E.coli SOLR"
/clone lib="OsHDAC1-overexpressing transgenic rice lambda
/clone lib="OsHDAC1-overexpressing transgenic rice lambda
phage cDNA library I (HDA1)"
/note="Vector: pBluescript SK(+); Site 1: EcoRI; Site 2:
XhoI; Callus was treated with ABA(20um) for lhour. cDNA
was inserted into lambda Uni-ZAP XR vector at 5' end with
EcoRI and 3' end with XhoI site. mRNA was derived from
/mol_type="mRNA"
/db_xrefs"taxon:9606"
/db_xrefs"taxon:9606"
/clone="IMAGE:1636750"
/clone_lib="NCI_CGAP_Kid3"
/note="Organ: kIdney; Vector: pT7T3D-Pac (Pharmacia) with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               rice Histone Deacetylase overexpression line.
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/cultivar="Nackdong"
/db_xref="taxon:39947"
                                                                                                                                            organism="Homo sapiens"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 6; he..
2.1e+06;
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BM400383/c
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19 b
1M0104A16R Mouse 10kb plasmid
clone UUGC1M0104A16 R, genomic
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Turkewitz,A.P., Karrer,K.M., Jahn,
Frankel,J. and Klobutcher,L.
EST from Tetrahymena thermophila,
Unpublished (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tetrahymena thermophila
Eukaryota; Alveolata; Ciliophora; Oligohymenophorea;
Hymenostomatida; Tetrahymenina; Tetrahymena.
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AZ361152
AZ361152.1 GI:10474852
GSS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Turkewitz AP
Molecular Genetics and Cell Biology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BM400383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Seq primer: T3
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920 E. 58th Street, Chicago,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tetrahymena thermophila
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: apturkew@midway.uchicago.edu
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                                                                                                                                                                                                                                                                                                                                                                                                                       preparation
Proc. Natl.
                                                                                                                                                                                                                                                                                                                                                                                                                  /db_xref="taxon:5911"
/clone lib="Chilcoat/Turkewitz cDNA (large fraction)"
/clone lib="Chilcoat/Turkewitz cDNA (large fraction)"
/note="Vector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    organism="Tetrahymena thermophila"
|mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /strain="CU428.1"
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100.0%; Pred. No. 2.2e+06;
tive 0; Mismatches 0;
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                                                                R, genomic survey sequence.
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                                                                                                                                                                                                                                                                                                                                  Score 11;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                             42 CCAGCCCATGC 52
    19 bp DNA linear GSS 14-DEC-200 (190506P02R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0506P02 R, genomic survey sequence.
AZ643124 AZ643124 GI:11770578 GSS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R. Mouse whole genome scaffolding with paired end reads from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                                                             11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0
Plate: 0104 row: A column: 16
Seq primer: CACACAGAAACAGCTATGACC
Class: plasmid ends
High quality sequence stop: 19.
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University of Utah
Rm. 308, Biomedical
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (bases 1 to 19)
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USA
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801 585 7177
                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligomucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNDA2 (gil4732114]gblAF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and neutrinal manuscript of the complementary to the insert adaptors and neutrinal manuscript.
                                                                                                                                                                                                                                                                                                                                                                                                                    purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XLIO-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    musculus C57BL/6J (male) was obtained from the Ja
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/).
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/clome_lib="Mouse 10kb plasmid UUGC1M library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /mol_type="genomic DNA"
/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                  Score 11;
Pred. No.
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1M0517123F Mouse 10kb plasmid clone UUGC1M0517123 F, genomic
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Insert Length: 10000 Std Error: 0.00
Plate: 0506 row: P column: 02
Seg primer: CACACAGGAAACAGCTATGACC
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.

Nouse, whole genome scaffolding with paired end reads from 10kb
                                          AZ648404.1 GI:11780837
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 High quality sequence stop: 19.
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801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gi 4732114 |gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent B. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/clone="UUGC1M0506P02"
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strain="C57BL/6J"
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Pred. No. 2.2e+06;
                                                                                                         genomic survey sequence.
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                               AZ995149
AZ995149.1
                                                                                               2M0280D22R Mouse 10kb plasmid clone UUGC2M0280D22 R, genomic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus

1 (bases 1 to 19)

1 (bases 1 to 19)

1 (bases 1 to 19)
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Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erry
Plate: 0517 row: I column: 2:
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20
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Contact: Robert B.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNava prepared from a derivative of pNava prepared from a derivative of pNava prepared from a derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       _tom/
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/mol_type="genomic DNA"
/strain="C57BL/6J"
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                                   GI:13866376
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Pred. No. 2.2e+06;
                                                                                                  genomic survey sequence.
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mid UUGC2M library Mus musculus
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          ACCESSION
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                                                                                                                                                                              RESULT 21
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Best Local
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22 bp mRNA linear EST 27-JUI or88e03.s1 NCI CGAP Lu5 Homo sapiens cDNA clone IMAGE:1602940 similar to TR:Q15816 Q15816 TRANSMEMBRANE PROTEIN JAGGED 1. [2 TR:O14902 ;contains TAR1.b2 MSR1 repetitive element ;, mRNA sequence.
AA989077
                                                                                                                                                                                                                                                                                                30 CACCTGCTGCT 40
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Insert Length: 10000 Std Error: C
Plate: 0280 row: D column: 22
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
High quality sequence stop: 19.
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Fax: 801 585 7177
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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Unpublished (2000)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus
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/clone_lib="Mouse 10kb plasmid UUGC2M library"
/note="Vector: pW042nv; Purified genomic DNA from.M.
musculus C57BL/6J (female) was obtained from the Jackson
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strain="C57BL/6J"
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Pred. No. 2.2e+06;
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Direct Submission
Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton,
                                                                    Hall, N., Bowman, S., Lennard, N.J., Doggett, J., Atkin, R., Chillingworth, C., Ormond, D., Harris, B., El-Sayed, N., Hou, L., Melville, S.E., Rajandream, M.A. and Barrell, B.G.
                                                                                                                                                                                                                                                                                                                                                                       T. brucei sheared genomic
                                                                                                                                                                                                                         Trypanosoma brucei
                                                                                                                                                                                                                                                                                                                       genomic survey sequence.
AL490359
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CDNA Library Preparation: M. Bento Soares, Ph.D. CDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html
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Insert Length: 760 Std Error: 0.00
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
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Unpublished (1997)
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National Cancer Institute, Cancer Genome Anat
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                                                                                                                                                                         Trypanosoma.
                                                                                                                                                                                             Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
                                                                                                                                                                                                                                                Trypanosoma brucei
                                                                                                                                                                                                                                                                                                AL490359.1 GI:11866449
                                                                                                                                                                                                                                                                                                                                                                                                TA312B10P
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                                                                                                                                                (bases 1 to 22)
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/rotee-"Organ: lung; Vector: pT773D-Pac (Pharmacia) with a modified polylinker; lst strand cDNA was prepared from neuroendocrine lung carcinoid, and was then primed with a Not I - oligo(dT) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library is normalized. Library was constructed by Bento Soares and M. Fatima Bonaldo. "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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/lab_host="DH10B"
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A clone
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POLYA=No.
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Guo, Y., Cai, Z. and Gan, S.
Transcriptome of Arabidopsis leaf senescence
Plant Cell Environ. 27 (5), 521-549 (2004)
Contact: Susheng Gan
Department of Horticulture
                                                                                                                                                                                                                                                                                                                                                                                                                  119 Plant Science, Cornell University, Ithaca, NY 14853-5904, USA Tel: 607 254 5418 Fax: 607 255 0599
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cornell University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CD531171
CD531171.1 GI:40451183
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23 bp
09L20 Arabidopsis Leaf Senescence
3', mRNA sequence.
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4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Bariell, Oxford University Press, 1999).
                                                                                                                                                                                                                                                                                                                                                                            Insert Length: 23
                                                                                                                                                                                                                                                                                                                                                                                                    Email: sg288@cornell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Arabidopsis thaliana
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Details of T. brucei sequencing at the Sanger Centre are available
at http://www.sanger.ac.uk/Projects/T_brucei/.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned population of Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically sheared
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nhl@sanger.ac.uk
Constructed at t
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/Glone lib="Arabidopsis Leaf Senescence Library" /note="Organ: Rosette Leaf; Vector: pBluscript SKII+; Site 1: EcoRI; Site 2: EcoRI; Senescent rosette leaves #5 and #6 (counted from the bottom) were harvested and immediately frozen in liquid N2. The leaves were visibly
                                                                                                                                                                                                                                       organism="Arabidopsis thaliana"
/mol_type="mRNA"
                                                                                                                                                                                                                                                                                                              location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /mol_type="genomic DNA"
/strain="TREU927"
                                                                                                                                                                    tissue_type="Leaf"
                                                                                                                                                                                              db_xref="taxon:3702"
                                                                                                                                                                                                               ecotype="Landsberg erecta"
                                                                                                                    dev stage="Yellow Leaf With Greenish Base Area" lab_host="E. coli"
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100.0%; Pred. No. 2.2e+06;
tive 0; Mismatches 0;
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Std Error:
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AZ331594/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (2000)
Contact: Robert B. Weiss
University of Utah
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10 GACTGCTACTG 20
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Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: C
Plate: 0059 row: D column: 12
Plate: 0059 row: D column: 12
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 24)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         High quality sequence stop: 24.
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AZ331594.1
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Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                             /lab_host="E. Coli strain XL10-Gold, T1-resistant, F /Clome_lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: PWA92nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jack
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /moi_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sex="Male
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100.0%; Pred. No.
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ORGANISM
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AI123486/c
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Best Local Similarity
Matches 11; Conserv
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome
Clone distribution: NCI-CGAP clone distribution
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 (bases 1 to 25)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi, Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     41143486 25 bp mRNA linear EST 01-OCT-1998 qa01e11.x1 Soares parathyroid tumor NbHPA Homo sapiens cDNA clone lMAGE.1683596 3' Similar to SW:BAT2_HUMAN P48634 LARGE PROLINE-RICH PROTEIN BAT2;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   High quality sequence stop: 1.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cDNA Library Preparation:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (1997)
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                                      Conservative
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                                                                                                                                         TTTTT-3'), double-stranded cDNA was size selected, ligated to Eco RI adapters (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of a modified pT7T3 vector (Pharmacia). Library went through one round of normalization to a Cot = 5. Library constructed by Bento Soares and M.Fatima Bonaldo. RNA from sporadic parathyroid adenomas was kindly provided by Dr. Stephen Marx, National Institute of Diabetes and Digestive and Kidney Diseases,
                                                                                                                                                                                                                                                                             /clone lib="Soares parathyroid tumor_NbHPA"
/note="Organ: parathyroid gland; Vector: pT7T3D
(Pharmacia) with a modified polylinker; Site 1:
Site_2: Eco RI; 1st strand cDNA was primed with
                                                                                                                                                                                                                                                                                                                                                                                                                       /tissue_type="parathyroid tumor"
/dev_stage="adult"
/lab_host="DH10B (ampicillin resistant)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        clone="IMAGE:1683596"
                                                    2.2%; Score 11; DB 1; Length 25; 100.0%; Pred. No. 2.2e+06;
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00.0%; Pred. No. 2.2e+06;
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                                    Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Source: IMAGE Consortium, LINL
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Trace considered overall poor quality
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 25)
Hilliar, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M.,
Holman, M., Hultman, M., Kucaba, T., Le, M., Lennon, G., Marra, M.,
Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F.,
Trevaskis, E., Waterston, R., Williamson, A., Wohldmann, P. and
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Yl24e12:r1 Soares breast 2NbHst Homo sapiens cDNA clone
IMAGE:159214 5' similar to gb:L19686 rnal MACROPHAGE MIGRATION
INHIBITORY FACTOR (HUMAN);, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fax:
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                                                                                                                                                                    Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: est@watson.wustl.edu
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Mammalia; Eutheria;
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Location/Qualifiers
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                                                                                                                                                                                                                                                         /mol_type="mRNA"
/db_xref="GDB:573305"
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                                                                                                                                                                  100.0%;
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                                                                                                                                                                  Score 11;
Pred. No.
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CG712540
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  survey sequence.
CG712540
CG712540.1 GI:37
GSS.
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yv08g12.rl Soares fetal liver spleen 1NFLS Homo sapiens cDNA clone
IMAGE:242182 5' similar to gb | M87933 | HUMAALUJ364 Human carcinoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi; Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 25)

1 (Dases 1 to 25)

Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M., Holman,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Pareaskis,E., Waterston,R., Williamson,A., Wohldmann,P. and
                                                                       CG712540
1119027E01.2EL_Y2 1119 -
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High quality sequence stops: 1
Source: IMAGE Consortium, LLNL
This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
Trace considered overall poor quality
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Contact: Wilson RK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: est@watson.wustl.edu
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                                                                                                                                                                                                       CCTGGGTGACA 5
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Location/Qualifiers
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314 286 1810
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                                                                                                                                                                                                                                                                                                                                                                               /mol_type="mRNA"
/db_xref="GDB:3791315"
/db_xref="taxon:9606"
/clone="IMAGE:242182"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                'sex="male"
                       GI:37738446
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o. 2.2e+06;
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Zea mays
                                                                                                                                                                                                                                                                              AU256278 3'-directed mouse cDNA BED0007961 3', mRNA sequence. AU256278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unpublished (2001)
Contact: Walbot V
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Generation of expressed sequence tags fro
Unpublished (2002)
Contact: Kikuya Kato
Graduate School of Biological Sciences
Nara Institute of Science and Technology
8916-5 Takayama, Ikoma, Nara 630-0101, Ja
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Department of Biological Sciences
Stanford University
855 California Ave, Palo Alto, CA
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Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD
clade; Panicoldeae; Andropogoneae; Zea.
                                                                                                                     1 (bases 1 to 26)
Kato, K. and Matoba, R.
                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                     Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Class: transposon-tagged.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: walbot@stanford.edu
Possible ligation site of ends cut by 2 different endonucleases.
Reverse complemented post-ligation sequence from source sequence
Plate: 1119027 row: E column: 01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tel: 650 723 2227
Fax: 650 725 8221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Maize genomic sequences found using engineered RescueMu transposon
                                                                                                                                                                                                                                                                 AU256278.1 GI:20319813
                                                                                                                                                                                                                          Mus musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /dev_stāge="adult"
/lab_host="DH10B"
/clone_lib="1119 - RescueMu Grid AA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /mol type="genomic DNA"
/cultivar="mixed background W23/A188/B73/K55"
/db_xref="taxon:4577"
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100.0%; Pred. No. 2.2e+06;
tive 0; Mismatches 0;
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  Japan
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JOURNAL COMMENT
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AUTHORS
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VERSION
KEYWORDS
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CF325420/c
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 305 AGAGCCTCGTG 315
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Fax: 81-743-72-5589
Email: ktatcobs aist-nara.ac.jp,
URL:http://love2.aist-nara.ac.jp/BED/index.html.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                Yongin, Kyeonggi, Ko
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H., Song,S.I., Kim,J.K., Kim,Y.-K. and Wahm,B.H. Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
Contact: Nahm B.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
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CF325420.1 GI:33799120
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                                 2.2%;
ilarity 100.0%;
Conservative (
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ilarity 100.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                   bhnahm@ggbio.com,
                                                                                                                                        'note="Vector: pBluescript SK(+); Site_1: EcoRI; Site_2: XhoI; cDNA was inserted into lamda Uni-ZAP XR vector at end with EcoRI and 3' end with NbOI site. mENA was prepared from Arabidopsis Jasmoinate Carboxyl
                                                                                                                                                                                                              /dev_stage="14 days after germination"
/lab_host="E.coli SOLM:
/clone_lib="AtJWT-overexpressing transgenic rice lambda
phage CDNA library (JWT1)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism="Mus musculus"
/mol_type="mRNA"
/db_xref="taxon:10090"
/clone="BED0007961"
                                                                                                                         methyltransferase overexpression
                                                                                                                                                                                                                                                                                                        /db_xref="taxon:39947"
/clone="JMT1--03-D02"
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/mol_type="mRNA"
/cultivar="Nackdong"
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/clone_lib="3'-directed mouse
                                                                                                                                                                                                                                                                                     tissue_type="leaf"
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                                                  Score 11; DB 6; L
Pred. No. 2.2e+06;
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                                     Mismatches
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ACCESSION
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                                                                                                                                                           RESULT 32
AZ318600/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Hall N
Pathogen Sequencing Unit
The Sanger Institute The Wellcome Trust Genome Cam
Hinxton, Cambridge, CB10 1SA, UK
Request for clones, please contact: Mike Lehane
Prof. M.J.Lehane
School of Biological Sciences,
University of Wales,
Bangor LL57 2UM
                                                                                                                                                                                                                                                                                      348 TTACAGAGTCA 358
                                                                                                                                                                                                                                                   11 TTACAGAGTCA 21
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                                                                                                                                                                                                                                                                                                                          11;
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                                                                                                                       AZ318600 29 ł
1M0037G14R Mouse 10kb plasmid
clone UUGC1M0037G14 R, genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lehane,M.J., Aksoy,S., Gibson,W., Kerhornou,A., Berriman,M., Hamilton,J., Soares,M.B., Bonaldo,M.F., Lehane,S. and Hall,N. Adult midgut expressed sequence tags from the tsetse fly Glossina morsitans morsitans and expression analysis of putative immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BX557759 Glossina morsitans morsitans adult infected gut Glossina morsitans morsitans cDNA clone Tse34e06_qlc, mRNA sequence.
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   All clones with suffix qlc are end of the cDNA all plc reads a the 3' end.
                                 Mus musculus
                                                    Mus musculus (house mouse)
                                                                                       AZ318600.1
                                                                                                          AZ318600
                                                                                                                                                                                                                                                                                                                      h 2.2%; Score 11; DB Similarity 100.0%; Pred. No. 2.: 11; Conservative 0; Mismatches
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1 (bases 1 to 29)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone="Tse34e06_q1c"
/tissue_type="adult infected gu
/clone_lib="Glossina morsitans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mol_type="mRNA"
/sub_species="morsitans"
/db_xref="taxon:37546"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             organism="Glossina morsitans"
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                                                                                       GI:10368531
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                                                                                                                         genomic survey sequence.
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Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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KEYWORDS
SOURCE
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                                                                1 1M0421B21R Mouse 10kb plasmid Uf
clone UUGC1M0421B21 R, genomic &
AZ602874
AZ602874.1 GI:11725064
GSS.
Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                       132 CCTGTACTGGG 142
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                                                                                                                                                                                                                                                                                                  28
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
Mouse whole genome scaffolding with paired end reads from 10kb
                                             Mus musculus
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Location/Qualifiers
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Insert Length: 10000 Std Error: 0.00
Plate: 0037 row: G column: 14
Seq primer: CACACAGGAAACAGCTATGACC
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Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              plasmid inserts
Unpublished (2000)
                                                                                                                                                                                                                                                                                                CCTGTACTGGG 18
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                                                              musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNA22 (gil 4732114 |gb|AF129072.1), a copy-number inducible derivative of plasmid Rl. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /mol_type="genomic DNA"
/strain="C57BL/6J"
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Pred. No. 2.3e+06;
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Best Local Similarity
Matches 11; Conserv
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Fax: 801 585 7177
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
                                                                Antirrhinum majus (snapdragon)
Antirrhinum majus
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University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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Insert Length: 10000 Std Error: (
Plate: 0421 row: B column: 21
Seq primer: CACACAGGAAACAGCTATGACC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pMD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-compotent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0421B21"
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100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                        Tel: 82 31 330 6193
Fax: 82 31 371
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Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnollophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University YongIn, Kyeonggi, Korea
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Le
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CF300184 30 bp mRNA linear EST:15-AUG-2003 7LEAF--04-H21.gl Rice leaf plasmid cDNA library II (7LEAF) Oryza sativa (japonica cultivar-group) cDNA clone 7LEAF--04-H21, mRNA
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Carl-von-Linne Weg 10, D-50829,
Location/Qualifiers
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Antirrhinum EST collection
Unpublished (2003)
Contact: Schwarz-Sommer Z
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Nahm B.H
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                                                                                                                                                                                                                                                                                                                                                                                  bhnahm@ggbio.com,
                                                                                                     /tissue_type="leaf"
/dev_stage="7 days after germination"
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/clone_lib="Rice_leaf_plasmid cDNA library II (7LBAF)"
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                                                                                                                                                                                                                                                                      /organism="Oryza sativa (japonica cultivar-group)"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "mol_type="mRNA"
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/db xref="taxon:4151"
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/clone_Tib="Antirrhinum majus whole plant"
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/clone="7LEAF--04-H21"
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    2.2%;
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    Score 11;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                           bhnahm@bio.myongji.ac.kr
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    DB 6; L,
2.3e+06;
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                       Length 30;
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            JMT1--03-N04.gl AtJMT-overexpressing transgenic rice lambda phage CDNA library (JMT1) Oryza sativa (japonica cultivar-group) cDNA clone JMT1--03-N04, mRNA sequence.
CF325654
CF325654.1 GI:33799587
                                                                                                                                                                                                                                                                                                                 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Arabidopsis Genome
Unpublished (2001)
Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R., Gadrinab,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L., Shinn,P., Zimmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Arabidopsis thaliana (thale cress)
Arabidopsis thaliana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SALK 118628.15.45.x Arabidopsis thaliana TDNA insertion lines Arabidopsis thaliana genomic clone SALK_118628.15.45.x, genom
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This is single pass sequence recovered from the left border of TDNA. This sequence lies within 300 bases of the 5' end of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ecker@salk.edu
                                                                                                                                                                                                                                                                                                                                                                                                              directly sequenced to determine the genomic sequence at the site of insertion. Details of the protocols used can be found at http://signal.salk.edu/tdna_protocols.html"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone lib="\bar{\rm A}rabidopsis thallana TDNA insertion lines" note="PCR was performed on Arabidopsis thallana lines each of which contains one or more TDNA insertion elements. The resultant fragment for each line was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /db_xref="taxon:3702"
/clone="SALK_118628.15.45.x"
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100.0%; Pred. No.
tive 0; Mismatc
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CL436292/c
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Email: hicksgg@cc.umanitoba.ca
U3NeoSV1 gene trap. Tag generated by plasmid rescue. Additional
u3NeoSV1 gene trap. Tag generated by plasmid rescue. Additional
sequence information and target gene cloning can be generated. ES
cell line harboring insertion mutation of target gene is available.
Sequence analysis available from
http://140.193.242.7/esdb/public_search_frame.php?PST=pST2678-NR.Se
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                                                                                                                                        Mammalian Functional Genomics Centre
Manitoba Institute of Cell Biology, Universitry of Manitoba
ON5029, 675 McDermot Ave, Winnipeg, MB R3E 0V9, Canada
Tel: 204 787 2133
Fax: 204 787 2190
                                                                                                                                                                                                                                           www.EScells.ca
Unpublished (2002)
Contact: Hicks GG
                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta;
Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae,
Ehrharroideae, Oryzeae, Oryza.
                                                                                                                                                                                                                                                                                                           Hicks, G.G
                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                               CL436292.1 GI:45570949
GSS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    genomic survey sequence.
CL436292
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
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larity 100.0%; Pred. No. 7.,
Conservative 0; Mismatches
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Location/Qualifiers
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/note="Vector: pBluescript SK(+); Site 1: EcoRI; Site 2: XhoI; cDNA was inserted into lamda Uni-ZAP XR vector at end with EcoRI and 3' end with XhoI site. mRNA was prepared from Arabidopsis Jasmoinate Carboxyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             methyltransferase overexpression line."
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[mol_type="maNA"
/cultivar="Nackdong"
/cultivar="Nackdong"
/db_xref="taxon:39947"
/clone="JMT1--03-N04"
/cissue_type="leaf"
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/lab_host="E.coli SOLR"
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o. 7.4e+06;
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Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 (bases 1 to 15)
Turkewitz,A.P., Karrer,K.M., Jahn,C., Orias,E., Kirk,K.E.,
Frankel,J. and Klobutcher,L.
EST from Tetrahymena thermophila, strain CU428.1, growing cells
Unpublished (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10;
                                                                                                                                                                                                                                                                                                                                                                                                               University of Chicago
920 E. 58th Street, Chicago, IL 60637,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tetrahymena thermophila
Eukaryota; Alveolata; Ciliophora; Oligohymenophorea;
Hymenostomatida; Tetrahymenina; Tetrahymena.
                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                   Seq primer: T3
                                                                                                                                                                                                                                                                                                                                                                                          Tel: 773 702 4374
Fax: 773 702 3172
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Turkewitz AP
Molecular Genetics and Cell Biology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BM397934 15 bp mRNA linear EST 17
5009-0-39-C05.t.l Chilcoat/Turkewitz cDNA (large fraction)
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Class: Gene Trap.
                                                                                                                                                                                                                                                                                                                                                                Email: apturkew@midway.uchicago.edu
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 ccecererce
                                  cccccicrce 91
                                                                       Conservative
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                                                                                                                                                      /db_xref="taxon:5911"
/clone lib="Chilcoat/Turkewitz cDNA (large fraction)"
/notee="Vector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
                                                                                                                                                                                                                                                   /organism="Tetrahymena thermophila"
/mol_type="mRNA"
/strain="CU428.1"
                                                                                                                                                                                                                                                                                                                                ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /cell type="Embryonic stem cell
/cell_line="D3H (J1 subclone)"
/clone_lib="MICB1"
/note="Vector: U3NeoSV1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /db xref="taxon:10090"
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/strain="129 sv"
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                                                                 2.0%; Score 10; DB 4; Lo
100.0%; Pred. No. 7.6e+06;
rative 0; Mismatches 0;
N
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                                                                                         SOURCE
ORGANISM
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (23-OCT-2003) Balzergue S., UMRGV, INRA/CNRS, 2 rue Gaston Cremieux, 91057 Evry cedex, FRANCE PCR was performed on DNA from transformants of Arabidopsis thaliana plants from INRA (Versailles). The DNA fragment(s) resulting from the PCR were directly sequenced from the left or the right border to determine the genomic sequence flanking the insertion. T-DNA derived sequences were removed. Information to order the corresponding mutant line and a link to a database providing a graphical display of the insertion site are available at http://dbsgap.versailles.inra.fr/publiclines/. This sequence has been generated in the framework of the French plant genomics program 'Genoplante' (http://www.genoplante.com and http://genoplante-info.infobiogen.ft).
                                                                                                                                                                                                                                                                                                                            289
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                                                                                                                                    BM401104
BM401104.1
                                                                                                                                                                                                                                                                                                                                                             10;
                                                                                                                                                                      16 bp ml
5009-0-82-F06.t.1 Chilcoat/Turkewitz
Tetrahymena thermophila cDNA, mRNA 86
                                                                  Tetrahymena thermophila
Eukaryota; Alveolata; Ciliophora;
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AJ590969.1 GI:37940593
GSS; left border; T-DNA flanking s
Arabidopsis thaliana (thale cress)
Arabidopsis thaliana
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     1 (bases 1 to 16)
Turkewitz, A.P., Karrer, K.M.,
Frankel, J. and Klobutcher, L.
                                                                                                        Tetrahymena thermophila
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Brunaud, V., Balzergue, S., Dubreucq, B., Aubourg, Chauvin, S., Bechtold, N., Cruaud, C., DeRose, R., Lepiniec, L., Caboche, M. and Lecharny, A. T-DNA integration into the Arabidopsis genome of pre-insertion sites EMBO Rep. 3 (12), 1152-1157 (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                       Hymenostomatida;
                                                                                                                                                                                                                                                                                                                                                                             Similarity
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-ucity 100.0%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Arabidopsis thaliana"

/mol type="genomic DNA"

/cultivar="Wassillewskija"

/db xrefs"taxon:3702"

/clone="577F01"
                                                                                                                                                                                                                                                                                                                                                                                                                                             /note="T-DNA flanking sequence
left border"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           clone_lib="Arabidopsis thaliana
                                                                                                                                          GI:18201157
                                                      Tetrahymenina;
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Pred.
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                    Jahn, C.,
                                                      Tetrahymena.
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7.6e+06;
hes 0;
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                                                                    Oligohymenophorea;
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                                                                                                                                                                                                         mRNA
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                                                                                                                                                                                  cDNA (large fraction)
                   Orias,E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Aubourg,S., Samson,F.,
Rose,R., Pelletier,G.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   genome depends on sequences
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ce, left border, clone
                                                                                                                                                                                                         linear
                   Kirk, K.E.,
                                                                                                                                                                                                       EST 17-JAN-2002
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BM397034/c
                                                                                                   ORIGIN
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Best Local
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                                                    Similarity
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BM397034 17 bp mRNA 1
5009-0-28-C07.t.2 Chilcoat/Turkewitz cDNA (1
Tetrahymena thermophila cDNA, mRNA sequence
                                                                                                                                                                                                                                                                                                                                                           University of Chicago
920 E. 58th Street, Cl
                                                                                                                                                                                                                                                                                                                                                                                               Contact: Turkewitz AP
Molecular Genetics and Cell Biology
                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (2002)
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BM397034
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Contact: Turkewitz AP
Molecular Genetics and Cell Biology
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Frankel,J. and Klobutcher,L.
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Eukaryota; Alveolata; Ciliophora;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           University of Chicago
920 E. 58th Street, Chicago,
                                                                                                                                                                                                                                                                                                        Email: apturkew@midway.uchicago.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hymenostomatida; Tetrahymenina; Tetrahymena.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tetrahymena thermophila
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773 702 3172
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773 702 3172
         Conservative
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                                                                                         /db xref="taxon:5911"
/clone lib="Chilcoat/Turkewitz cDNA (large fraction)"
/notee="Vector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (20
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
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/note="Vector: BlueScript2 SK; Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
                                                                                                                                                                                              /mol_type="mRNA"
/strain="CU428.1"
                                                                                                                                                                                                                                                                   Location/Qualifiers
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2.0%; bu
100.0%; Pr
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100.0%; Pred. No.
tive 0; Mismatcl
                                                                                                                                                                                                                                                                                                                                                               Chicago, IL 60637, USA
       Score 10; DB; Pred. No. 7.6
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3. 7.6e+06;
0;
                         DB 4; Le
7.6e+06;
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                                        Length 17;
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RESULT 43
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EST from Tetrahymena thermophila, strain CU428.1, growing cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tetrahymena thermophila
Eukaryota; Alveolata; Ciliophora; Oligohymenophorea;
Hymenostomatida; Tetrahymenina; Tetrahymena.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BM397514 17 bp mRNA linear EST 17 5009-0-33-H05.t.1 Chilcoat/Turkewitz cDNA (large fraction)
                                                                                                                                                                                                                     Tetrahymena thermophila
Tetrahymena thermophila
Eukaryota; Alveolata; Ciliophora; Oligohymenophorea;
Contact: Turkewitz AP
Molecular Genetics and Cell Biology
University of Chicago
920 E. 58th Street, Chicago, IL 606
Tel: 773 702 4374
Fax: 773 702 3172
                                                                                                          Unpublished (2002)
                                                                                                                                                Turkewitz, A.P., Karrer, K.M., Jahn, C., Orias, E., Kirk, K.E. Frankel, J. and Klobutcher, L.
                                                                                                                                                                                                                                                                                               BM398743
BM398743.1 GI:18198796
                                                                                                                                                                                                                                                                                                                                18 bp mRNA 15009-0-49-F05.t.1 Chilcoat/Turkewitz cDNA (1701-2014) mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          University of Chicago
920 E. 58th Street, Chicago, IL 60637,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Molecular Genetics and Cell Biology
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                                                                                                                               EST from Tetrahymena thermophila,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.0%; Score 10; DB 4; Le Similarity 100.0%; Pred. No. 7,6e+06; O: Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: apturkew@midway.uchicago.edu
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                                                                                                                                                                                                      Hymenostomatida; Tetrahymenina;
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773 702 3172
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/clone lib="Chilcoat/Turkewitz cDNA (large fraction)"
/note="Vector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
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/strain="CU428.1"
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                                      IL 60637,
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                                                                                                                             strain CU428.1, growing cells
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BQ590149
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VERSION
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                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Weisshaar B
ADIS DNA core facility at MPIZ
Max-Planck-Institute for Plant Breeding Research
Carl-von-Line Weg 10, 50829 Koeln, Germany
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Construction of a 'unigene' cDNA clone set by oligonucleotide fingerprinting allows access to 25 000 potential sugar beet genes Plant J. 32 (5), 845-857 (2002)
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18 bp mRNA linear EST 06-DEC-
E012845-024-019-B19-T7 MPIZ-ADIS-024-storage root Beta vulgaris
CDNA Clone 024-019-B19 3-PRIME, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M., 
Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Lehrach,H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta, Spermatophyta, Magnoliophyta, eudicotyledons, core eudicots, Caryophyllales, Amaranthaceae, Beta.

1 (bases 1 to 18)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fax: 00492215062851
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/clone lib="chilcoat/Turkewitz cDNA (large fraction)"
/note="Vector: BlueScript2 SK+, Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
SP6-SalI-CCACGCGTCCG-5prime-cDNA-polyA-CC-NotI-T7; Sequencing granted in the context of the GABI-Beet
                                                                                   /clone lib="MPIZ-ADIS-024-storage root" /clone lib="MPIZ-ADIS-024-storage root" /note="Vector: pCMVSPORT6; Site 1: Sall; Site 2: NotI; /cDNA library from sugar beet, library provided by KWS Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact
                                                               b.schulz@kws.de; cloning sites Sall-NotI, primer sites and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        organism="Tetrahymena thermophila"
/mol_type="mRNA"
                                                                                                                                                                                                                /db_xref="GABI:189975"
/db_xref="taxon:161934"
/clone="024-019-B19"
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                                                                                                                                                                                                                                                                                                  /organism="Beta vulgaris"
/mol_type="mRNA"
/cultivar="KW82320 (double haploid, monogerm breeding
                                                                                                                                                                      /tissue_type="storage root"
/lab_host="EMDH10B"
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AI049374/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19 bp mRNA linear EST 08-JUL-: ub33a03.rl Soares_thymus_2NbMT Mus musculus cDNA clone IMAGE:1379500 5' similar to SW:KDGZ_HUMAN Q13574 DIACYLGLYCEROL KINASE, ZETA ;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10;
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                                                                                                                                                                                                                                                                                                                                          Trace considered overall poor quality
Possible reversed clone: similarity on wrong strand
Seq primer: -28ml3 rev2 ET from Amersham
High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                   Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
TMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Washington University School of MedicineP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Marra M/Mouse EST Project WashU-HHMI Mouse EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Marra,M., Hillier,L., Allen,M., Bowles,M., Dietrich,N., Dubu, Geisel,S., Kucaba,T., Lacy,M., Le,M., Martin,J., Morris,M., Schellenberg,K., Steptoe,M., Tan,F., Underwood,K., Moore,B., Theising,B., Wylie,T., Lennon,G., Soares,B., Wilson,R. and Waterston,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi, Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Mus.
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EST.
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       project, local PI: Dr. Katharina Schneider, coordinator: Prof. Christian Jung; Sequence submission managed by RZPD/GABI-Primary database: http://gabi.rzpd.de"
                                                                                                                                                                         /tissue_type="Thymus"
/dev_stage="4 weeks"
/lab_host="DH108"
                                                                                                                                                                                                                                                  /organism="Mus musculus"
/mol_type="mRNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
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                                                                                                                                                                                                                                       clone="IMAGE:1379500"
                                                                                                                                                                                                                       sex="male"
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BM395679/c
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150 CCGGACCGGC 159
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                                                                                                  1 (bases 1 to 19)
Turkewitz,A.P., Karrer,K.M., Jahn,
Frankel,J. and Klobutcher,L.
BST from Tetrahymena thermophila,
Unpublished (2002)
                                                                                                                                                                                          Tetrahymena thermophila
Tetrahymena thermophila
Eukaryota, Alveolata; Ciliophora; Oligohymenophorea;
Hymenostomatida; Tetrahymenina; Tetrahymena.
                                                Molecular Genetics and Cell Biology
University of Chicago
                                                                                                                                                                                                                                                                                                                   5009-0-10-A05.t.1 Chilcoat/Turkewitz cDNA (large fraction) Tetrahymena thermophila cDNA, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Blum, H., Bauersachs, S.,
EST (Blum, et al.)
Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa;
Mammalia; Eutheria;
1 (bases 1 to 19)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ALO42746 19 bp mRNA linear DKFZp434C1822_r1 434 (synonym: htes3) Homo sapiens DKFZp434C1822, mRNA sequence.
ALO42746
                                                                                       Contact: Turkewitz AP
                                                                                                                                                                                                                                                                                    BM395679.1 GI:18195732
                920 E. 58th Street,
Tel: 773 702 4374
                                                                                                                                                                                                                                                                                                        BM395679
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    Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /dev_stāge="adult"
/lab_host="DH10B"
/clone_lib="434 (synonym: htes3)"
/note="Vector: pSport1; Site_1: Not1; Site_2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /clone="DKFZp434C1822"
/tissue_type="testis"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    organism="Homo sapiens"
(mol_type="mRNA"
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100.0%; Pred. No.
                                                                                                                                                          Karrer, K.M., Jahn, C.,
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Primates;
                               Chicago,
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                                                                                                                       strain CU428.1, growing cells
                                                                                                                                                                                                                                                                                                                                                          mRNA
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                                                                                                                                                          Orias, E., Kirk, K.E.,
                                   USA
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Seq pr
 34 TGCTGCTTCC 43
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13104 Arabidopsis Leaf Senescence Library Arabidopsis thaliana cDNA
37, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: sg288@cornell.edu
Insert Length: 19 Std |
Seg_primer: T7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   119 Plant Science, Cornell University, Tel: 607 254 5418 Fax: 607 255 0599
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Guo,Y., Cai,Z. and Gan,S.
Transcriptome of Arabidopsis leaf senescence
Plant Cell Environ. 27 (5), 521-549 (2004)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Arabidopsis thaliana (thale cress)
Arabidopsis thaliana
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CD532073.1 GI:40452085
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Department of Horticulture
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Susheng Gan
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                                     Conservative
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                                                                                                                              /dev_stage="Yellow Leaf With Greenish Base Area"
//ab host="E, coli"
/clone_lib="Arabidopsis Leaf Senescence Library"
/note="Organ: Rosette Leaf; Vector: pBluscript SKII+;
/site_1: EcoRI, Site_2: EcoRI; Senescent rosette leaves #5
and #6 (counted from the bottom) were harvested and
amediately frozen in liquid N2. The leaves were visibly
yellow excepted for the leaf base areas that were still
greenish. "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /db xref="taxon:5911"
/clone lib="Chilcoat/Turkewitz cDNA (large fraction)"
/clone lib="Chilcoat/Turkewitz cDNA (large fraction)"
/note="Vector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
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/mol_type="mRNA"
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                                                                                                                                                                                                                                                                                                                                                              organism="Arabidopsis thaliana"
|mol_type="mRNA"
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                                                                                                                                                                                                                                                                                                                                              ecotype="Landsberg erecta"
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7.7e+06;
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AUTHORS
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Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., Le Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H. Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CF305339 19 bp mRNA linear EST 15-AUG-2003 CLD1--01-H03.bl Rice cold treated leaf plasmid cDNA library (CLD1) Oryza sativa (japonica cultivar-group) cDNA clone CLD1--01-H03, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Nahm B.H.
Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
   Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: bhnahm@ggbio.com, bhn
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Yongin, Kyeonggi, Ko
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae; Oryzeae; Oryza.
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CF305339.1 GI:33677100
                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 19)
                                                                                                                                                                                            AZ314110
AZ314110.1 GI:10359675
                                                                                                                                                                                                                             1M0030E16R Mouse 10kb plasmid to clone UUGC1M0030E16 R, genomic
                                                                                                                                                                                                                                                                         AZ314110
                                                                                                                                      Mus musculus
                                                                                                                                                    Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
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                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /tissue_type="leaf"
/dev_stage="14 days after germination"
/lab_host="B.coli DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /mol_type="mRNA"
/cultivar="Nackdong"
/db_xref="taxon:39947"
/clone="CLD1--01-H03"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            organism="Oryza sativa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Clone lib="Rice cold treated leaf plasmid cDNA library
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Pred. No.
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                                                                                                                                                                                                                                 19 bp DNA linear GSS 29 smid UUGC1M library Mus musculus nomic survey sequence
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                                                           Duval, B., Hamil, C.,
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AZ394490
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19 bp DNA linear GSS 03-OCT-200
1M0158005F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0158005 F, genomic survey sequence.
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (Dases 1 to 19)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., 181am,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R.
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GSS.
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Location/Qualifiers
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Insert Length: 10000 Std Error: 0.00
Plate: 0030 row: E column: 16
Seq primer: CACACAGGAAACAGCTATGACC
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801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gil 4732114 |gb| AFT29072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/strain="C57BL/6J"
/db_xref="taxon:10090"
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Pred. No. 7.7e+06;
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Insert Length: 10000 Std Error: 0
Plate: 0158 row: O column: 05
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
High quality sequence stop: 19.
Location/Qualifiers
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                 Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von
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                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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Contact: Robert B.
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Fax: 801 585 7177
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/strain="C57BL/6J"
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100.0%; Pr
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Insert Length: 1000 Std Error: 0.00
Plate: 0306 row: J column: 12
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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Fax: 801 585 7177
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                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 19)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Laboratory Mouse DNA Resource
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Bukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleosto Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; 1 (bases 1 to 19)

Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weise, R.
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19 bp DNA linear GSS 13-DEC-20.
1M0467A14F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0467A14 F, genomic survey sequence.
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Location/Qualifiers
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Insert Length: 10000 Std Err
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Fax: 801 585 7177
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/mol_type="genomic DNA"
/strain="C57BL/6J"
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 19)

Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                       19 bp DNA linear GSS 14-DEC-20
1M0513016R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0513016 R, genomic survey secuence
AZ647364
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Location/Qualifiers
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Insert Length: 10000 Std Error:
Plate: 0467 row: A column: 14
Seq primer: CGTTGTAAAACGACGGCCAGT
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/strain="C57BL/6J"
/db_xref="taxon:10090"
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Pred. No. 7.7e+06;
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2M0164104F Mouse 10kb plasmid clone UUGC2M0164104 F, genomic
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Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
High quality sequence stop: 19.
Location/Qualifiers
            1 (bases 1 to 19)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T.,
Reilly,M., Rose,M., Rose,M., Stokes,R., Tingey,A., von
                                                                                                                                                                   Mus musculus
    Niederhausern, A. and Wright, D., Weiss, R.
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Insert Length: 10000 Std Error: 0.00
Plate: 0513 row: O column: 16
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Mammalia; Eutheria; Rodentia;
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strain="C57BL/6J"
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    Morio, T., Urushihara, H., Saito, T., Ugawa, Y., Mizuno, H., Yos
Yoshino, R., Mitra, B.N., Pi, M., Sato, T., Takemoto, K., Yasuka
Williams, J., Maeda, M., Takeuchi, I., Ochiai, H. and Tanaka, Y.
Developmental cDNA in Dictyostelium discoideum
Unpublished (1998)
                                                                                                                  Dictyostelium discoideum
Dictyostelium discoideum
Eukaryota; Mycetozoa; Dictyosteliida; Dictyostelium.
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Insert Length: 10000 Std Erro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pw942 (gi 4732114 |gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-elected for a 9.5 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/Clone_lib="Mouse 10kb plasmid UUGCIM library"/note="Vector: PW042nv, Purified genomic DNA from M.musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Biomedical Polymers Research Bldg.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mol_type="genomic DNA"
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; Pred. No. 7.7e+06;
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                                                                                 Yoshida, M.,
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BX551623
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Glossina morsitans morsitans
Glossina morsitans morsitans
Eukaryota, Metazoa, Arthropoda, Hexapoda, Inse
Neoptera, Endopterygota, Diptera, Brachycera,
Hippoboscoidea, Glossinidae, Glossina.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BX551623 Glossina morsitans morsitans adult infected gut Glossina morsitans morsitans adult infected gut Glossina morsitans morsitans cDNA clone Tsell9f08_qlc, mRNA sequence.

BX551623
                                                                                                                                                                                                                        All clones with suf end of the cDNA all the 3' end.
                                                                                                                                                                                                                                                                                Prof. M.J.Lehane
School of Biological Sciences,
University of Wales,
Bangor LL57 2UW
                                                                                                                                                                                                                                                                                                                                                          The Sanger Institute The Wellcome Trust Genome Campus Hinxton, Cambridge, CB10 1SA, UK
Request for clones, please contact: Mike Lehane
                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Hall N
Pathogen Sequencing Unit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lehane,M.J., Aksoy,S., Gibson,W., Kerhornou,A., Berriman,M., Hamilton,J., Soares,M.B., Bonaldo,M.F., Lehane,S. and Hall,N. Adult midgut expressed sequence tags from the testse fly Glossina morsitans morsitans and expression analysis of putative immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      response genes
Genome Biol. 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14519198
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: hideko@biol.tsukuba.ac.jp
PROJECT = Dictyostelium discoideum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Institute of E
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Hideko Urushihara
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
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                      /db_xref="taxon:37546"
/clone="Tsel19f08 q1c"
/tissue_type="adult infected gut"
/clone_lib="Glossina morsitans morsitans adult infected gut"
/note="country: Zimbabwe; EST from adult gut infected with
                                                                                                                                  /organism="Glossina morsitans morsitans"
/mol type="mRNA"
/____
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Dictyostelium
/mol_type="mRNA"
/strain="AX4"
                                                                                                                    sub_species="morsitans"
                                                                                                                                                                                                     location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /dev_stage="slug"
/clone_lib="Dictyostelium discoideum SL (H.Urushihara)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /db_xref="taxon:44689"
/clone="SLK222"
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/ 100.0%; Pr
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Pred. No.
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ACCESSION
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CF307503/c
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Matches 1
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                          10;
AZ303903 20 bp DNA linear GSS 29 1M0003B18R Mouse 10kb plasmid UUGC1M library Mus musculus clone UUGC1M0003B18 R, genomic survey sequence. AZ303903 AZ303903.1 GI:10339339
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeonggi, Korea Tel: 82 31 330 6193 Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 (bases 1 to 20)
Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H., Shin,Y.C.,
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CF307503.1 GI:33679264
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                              /dev_stage="proliferated callus on 2N6 media for 2 weeks"
/lab host="E.coli SOLR"
/clome lib="05HDAC1-overexpressing transgenic rice lambda
/clome lib="05HDAC1-overexpressing transgenic rice lambda
phage cDNA library I (HDA1)"
/note="Vector: pBluescript SK(+); Site 1: EcoRI; Site 2:
/note="Vector: pBluescript SK(+); Site 1: EcoRI; Site 2:
/note="Vector: pBluescript SK(+); For lhour. cDNA
was inserted into lambda Uni-ZAP XR vector at 5' end with
was inserted into lambda Uni-ZAP XR vector at 5' end with
EcoRI and 3' end with XhoI site. mRNA was derived from
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/mol_type="mRNA"
/cultivar="Mackdong"
/db_xref="taxon:39947"
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100.0%; Pred. No. 7.7e+06;
tive 0; Mismatches 0;
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.00.0%; Pred. No. 7.7e+06; 
.ve 0; Mismatches 0; Indels
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FAX: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 1000 Std Error: 0.4
Plate: 003 row: B column: 18
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
High quality sequence stop: 20.
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                      AZ307491 20 bp DNA linear GSS 29-SEP-20-
1M0009C13F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0009C13 F, genomic survey sequence.
AZ307491
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AZ307491.1 GI:10346544
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished (2000)
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /mol_type="genomic DNA"
/strain="C57BL/6J"
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Mammalia; Eutheria; Rodentia;
1 (bases 1 to 20)
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Mus musculus (house mouse)
Mus musculus
AZ407675 DD DWA TIME

1M0178E04R Mouse 10kb plasmid UUGCIM library Mu

clone UUGCIM0178E04 R, genomic survey sequence.

AZ407675

AZ407675.1 GI:10531688
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Class: plasmid ends
High quality sequence stop: 20.
Location/Qualifiers
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Insert Length: 1000 Std Erro
Plate: 0009 row: C column: 13
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Fax: 801 585 7177
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University of Utah
Genome Center
University of Utah
Rm. 308, Biomedical Polymers Res
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/strain="C57BL/6J"
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Query Match 2.0%; Score 10; DB 8; L
Best Local Similarity 100.0%; Pred. No. 7.7e+06;
Matches 10; Conservative 0; Mismatches 0;
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Mus musculus (house mouse)
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Fax: 801 585 7177

Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0178 row: E column: 04
Seg primer: CACACAGGAAACAGCTATGACC
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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High quality sequence stop: 20.
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                                                          1M0311C24F Mouse 10kb plasmid UUGC1M library Muclone UUGC1M0311C24 F, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0178E04"
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/mol_type="genomic DNA"
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GI:10649799
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AZ659783/c
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AZ659783 20 bp DNA linear GSS 14 1M0537M23F Mouse 10kb plasmid UUGC1M library Mus musculus clone UUGC1M0537M23 F, genomic survey sequence. AZ659783 GI:11796929
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0311 row: C column: 24
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
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Fax: 801 585 7177
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Mus musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
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/strain="C57BL/6J"
/db_xref="taxon:10090"
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                                                                                                                                    AZ809111/c
                                                                                                                                                                RESULT 66
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AZ809111 20 bp DNA linear 2M0072P23R Mouse 10kb plasmid UUGC1M library Mus r clone UUGC2M0072P23 R, genomic survey sequence. AZ809111 GI:12975140
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Fax: 801 585 7177

Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0537 row: M column: 23
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
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University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Rese
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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/strain="C57BL/6J"
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                                                                AZ835099 20 bp DNA linear GSS 20-FEB-200 2M0129107F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC2M0129107 F, genomic survey sequence.
                         AZ835099.1 GI:13005007
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Fax: 801 585 7177

Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00

Plate: 0072 row: P column: 23
Seq primer: CACACAGGAAACAGCTATGACC
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Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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Mus musculus (house mouse)
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/mol_type="genomic DNA"
/strain="C57BL/6J"
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/clone="UUGC2M0072P23"
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Seq primer: CGTTGTAAAACGACGGCCAGT
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Insert Length: 10000 Std Error:
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Fax: 801 585 7177
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Mus musculus (house mouse)

Mus musculus
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/strain="C57BL/6J"
/db_xref="taxon:10090"
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GSS 06-MAR-2004 survey

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Gaps

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Submitted (07-JAN-2002) Hong-Seog Park, Korea Research Institute of Submitted (07-JAN-2002) Hong-Seog Park, Korea Research Institute of Bioscience and Biotechnology (KRIBB), Genome Research Center (GRC), 52, Oun-dong, Yusong-gu, Daejeon 305-333, Korea (E-mail:redstone@mail.kribb.re.kr, URL:http://phs.grc.kribb.re.kr/,
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                                                                                             2 k. Dases 1 to 20)
2 k. Dases 1 to 20)
Park, H., Kim, Y., Kim, S., Han, Y., Woo, T., Park, K., Eun, C.J.,
Hoon, S.T., Chu, M., Kim, H., Joo, S., Kim, C., Song, W. and Yoo, H.
Direct Submission
                                                                                                                                                                                                   Park,H., Kim,Y., Kim,S., Han,Y., Woo,T., Park,K., Eun,C.J., Hoon,S.T., Chu,M., Kim,H., Joo,S., Kim,C., Song,W. and Yoo,H. BAC end sequences of Library RP-43
                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Pan.
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AG191409.1 GI:45223585
                                                                                                                                                                                                                                                                                                                                                                                                                                                          AG191409 20 bp DNA Pan troglodytes DNA, clone: RP43-067F16.TJ,
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Clones are derived from the chimpanzee BAC library RP-43 This BAC end was generated during the R&D process and may have higher chance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (07-JAN-2002) Hong-Seog Park, Korea Research Institute of Bioscience and Biotechnology (KRIBB), Genome Research Center (GRC); 52, Oun-dong, Yusong-gu, Daejeon 305-333, Korea (E-mail:redstone@mail.kribb.re.kr, URL:http://phs.grc.kribb.re.kr/,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Park,H., Kim,Y., Kim,S., Han,Y., Woo,T., Park,K., Eun,C.J., Hoon,S.T., Chu,M., Kim,H., Joo,S., Kim,C., Song,W. and Yoo,H. Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Park, H., Kim, Y., Kim, S., Han, Y., Woo, T., Park, K., Eun, C.J., Hoon, S.T., Chu, M., Kim, H., Joo, S., Kim, C., Song, W. and Yoo, H. BAC end sequences of Library RP-43
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Pan troglodytes
Eukaryota; Metazoa; Chordata
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Pan.
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/clone_lib="RP-43 Chimpanzee Male
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/db_xref="taxon:9598"
/clone="RP43-061M08.T7"
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100.0%; Pred. No. 7.7e+06;
vative 0; Mismatches 0;
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Submitted (07-JAN-2002) Hong-Seog Park, Korea Research Institute of Bioscience and Biotechnology (KRIBB), Genome Research Center (GRC); 52, Oun-dong, Yusong-gu, Daejeon 305-333, Korea (E-mail:redetone@mail.kribb.re.kr, URL:http://phs.grc.kribb.re.kr/,
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Tel:82-42-866-7181, Fax:82-42-860-4409)
Clones are derived from the chimpanzee BAC library RP-43 This BAC
Clones are derived from the R&D process and may have higher chan
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R.Site 1 : EcoRI
R.Site 2 : EcoRI.
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    /cell_type="lymphocytes"
                                                                    /mol_type="genomic DNA"
/db_xref="taxon:9598"
                                                                                                                                                       location/Qualifiers
                                                clone="RP43-077L17.TJ"
                                                                                                             organism="Pan troglodytes"
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chance

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77 AGGACCCGCG 86
                                                                                        Park,H., Kim,Y., Kim,S., Han,Y., Woo,T., Park,K., Eun,C.J., Hoon,S.T., Chu,M., Kim,H., Joo,S., Kim,C., Song,W. and Yoo,H. BAC end sequences of Library RP-43
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Clones are derived from the chimpanzee BAC library RP-43 This BAC end was generated during the R&D process and may have higher chance of clone tracking errors.
Park,H., Kim,Y., Kim,S., Han,Y., Woo,T., Park,K., Eun,C.J., Hoon,S.T., Chu,M., Kim,H., Joo,S., Kim,C., Song,W. and Yoo,H.
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Mammalia; Eutheria; Primates;
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Similarity 100.0%; P
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R.Site 1 : EcoRI
R.Site 2 : EcoRI.
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/clone_lib="RP-43 Chimpanzee Male BAC Library"
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              AU013419 21 bp mRNA linear EST 03-AU AU013419 Schizosaccharomyces pombe late log phase cDNA Schizosaccharomyces pombe cDNA clone spc08126, mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tel:82-42-866-7181, Fax:82-42-860-4409)
Clones are derived from the chimpanzee BAC library RP-43 This BAC end was generated during the R&D process and may have higher chance
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (07-JAN-2002) Hong-Seog Park, Korea Research Institute of Bioscience and Biotechnology (KRIBB), Genome Research Center (GRC); Coun-dong, Yusong-gu, Daejeon 305-333, Korea (E-mail:redstone@mail:kribb.re.kr, URL;http://phs.grc.kribb.re.kr/,
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Pan.
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'clone="RP43-084C17.T7"
                                                                                                                                                                                                                                                                                                                                                                                                                  organism="Pan troglodytes"
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>_lib="RP-43 Chimpanzee Male BAC Library"
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;; Pred. No. 7.7e+06;
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Pred. No.
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AU013601/c
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Genome Research Group
National Institute of Radiological Sciences
National Institute of Inage-ku, Chiba, Chiba
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National Institute of Radiological Sciences
National Institute of Radiological Sciences
National Institute of Radiological Sciences
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Eukaryota; Fungi; Ascomycota; Schizosaccharomycetes;
Schizosaccharomycetales; Schizosaccharomycetaceae;
                                                                                                     9-1, Anagawa-4-chome, Inage-ku,
Email: morimyo@nirs.go.jp.
Location/Qualifiers
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1 (bases 1 to 21)
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Morimyo,M. and Mita,K.
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Location/Qualifiers
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into the Smal site of Ml3mp19 DNA and the direction of DNA
sequences was not always from 5' to 3'. The cDNA data of
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Contact: Mitsuoki Morimyo
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l: morimyo@nirs.go.jp.
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AC AAF30374;

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DT 14-MAY-2001 (first entry)

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DT 01 14-MAY-2001 (first entry)

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Checkpoint gene chfr 3'

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KW Checkpoint with forkhead asso
Checkpoint with forkhead associated domain and ring finger; Chfr; human; mitosis; cell cycle; tumour; diagnosis; antitumour; drug screening;
                                                           Human checkpoint gene chfr 3' PCR primer.
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AAS449421

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Abq72778 Insulin r
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Aah26573 Mouse addi
Acd06338 Forward R
Adm16634 Human NOV
Az294779 AIP bindi
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Abn12749 Human GDM
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Abn12747 Human GDM
Abn12740 Human GDM
Abn12750 Human GDM
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Aas84451 SPINKS 5'
Abk49942 PCR prime
Abk66679 Human gen
Aad35046 Zebrafish
Acf30912 Rice chro
Abx99708 A. 11quef
Adi09971 Rice chro
Ado47999 Human HIP
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Human
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AAF30360/c
ID AAF30360 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 18
                                                                                                                                                          Checkpoint with forkhead associated domain and ring mitosis; cell cycle; tumour; diagnosis; antitumour; ubiquitin-protein ligase; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel nucleic acid sequence of mitotic checkpoint gene encoding a checkpoint with forkhead-associated domain and ring finger protein, diagnosing tumorigenic cells and in screening for anticancer drugs.
    14-JUN-2000; 2000WO-US016391.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18 BP; 8 A; 6 C; 3 G; 1 T; 0 U; 0 Other;
                                             08-FEB-2001
                                                                                                                                                                                                                                            Human
                                                                                                                                                                                                                                                                                 14-MAY-2001
                                                                                                                                                                                                                                                                                                                        AAF30360,
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                                                                                                                         Homo sapiens.
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                                                                                                                                                                                                                                          checkpoint
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18;
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                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                        gene chfr 3'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3.6%;
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Pred. No.
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104 105 107 108

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95 96 97 98 99 100 101 102 103

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PCR

primer

finger; Chfr; human;
drug screening;

Mismatches

0

Gaps

0

DB 5; 1.2e+02;

Length 18;

therapies

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ARESULT 3
AAF30364/c
ID AAF303
XX AAF303
AC AAF303
AC Human
YX Checkp
KW Mitosi
KW ubiqui
XX Homo 6
YX WO2001
YX WO3001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence is that of a 3' PCR primer, used with the 5' primer CC given in AAF30359, to amplify a cDNA fragment corresponding to nucleotides 904-1753 of the human chfr sequence given in AAF30352. The CC chfr gene encodes the human mitotic checkpoint protein Chfr (see CC AAB20219), which is required for regulation of the transition of cells (from prophase to metaphase during mitosis. Loss of expression of Chfr is associated with a predisposition to tumourigenesis upon exposure to cmitotic stress. A set of primers (see AAF3033-76) was used to amplify regions spanning the entire chfr coding region in order to determine whether the chfr gene is mutated in any of the human cancer cell lines SW480, DLD1, HT29, HCT116, SAOS2, U2OS, IMR5 and NGP. A mutation leading to a Val-580 to Met amino acid substitution was identified in the chfr gene of U2OS cells. Chfr polypeptides and chfr mucleic acids are used in methods of diagnosing tumourigenic cells and of screening for drugs which can inhibit the activity of Chfr in a cancer cell, rendering it more constituted to additional antitumour therapies
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                       Checkpoint with forkhead associated domain and ring finger; Chfr; human; mitosis; cell cycle; tumour; diagnosis; antitumour; drug screening; ubiquitin-protein ligase; PCR primer; ss.
                                                                                                                                                                                                                     14-JUN-2000; 2000WO-US016391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human checkpoint
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-MAY-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAF30364 standard; DNA; 18
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                           WPI; 2001-182927/18
                                                                        Halazonetis T,
                                                                                                                                                                                                                                                                                                                  WO200109150-A2
                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                        29-JUL-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3; Page 38; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CGTTTTGTGAGCTCAACC 196
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                                                                                                                        INST ANATOMY & BIOLOGY.
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                                                                           Scolnick D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Scolnick D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            gene chfr 3' PCR primer.
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                                           Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel nucleic acid sequence of mitotic checkpoint gene encoding a checkpoint with forkhead-associated domain and ring finger protein diagnosing tumorigenic cells and in screening for anticancer drugs
                                                                                                                              Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 3; Page 38; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        present sequence is that of a 3' PCR primer, used with the 5'
                                                                 Similarity
                                                                                                                            BP; 8 A;
                                             Conservative
                                                               3.6%;
                                                                                                                              6 C; 3
                                       ; Score 18; DB 5; Leix
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                                                                                                                                Other;
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                                                                                  Length 18;
                                             Indels
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328
18
                  GGAGTGTTTCTGCTGTCT 345
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RESULT 4
AAF30366/c
ID AAF303
XX AAF303
XX AAF303
XX Checkp
DT 14-MAY
XX Checkp
KW mitcsi
KW mitcsi
KW ubiqui
XX Homo 8
XX HOMO 9
XX HOMO 9
XX HOMO 9
XX HOMO 9
XX HOMO 14-JUN
XX HOM AAF30366; Human checkpoint gene chfr 3' PCR primer. 14-MAY-2001 AAF30366 standard; DNA; 18 (first entry)

Checkpoint with forkhead associated domain and ring finger; Chfr; human; mitosis; cell cycle; tumour; diagnosis; antitumour; drug screening; ubiquitin-protein ligase; PCR primer; ss.

WO200109150-A2

08-FEB-2001.

14-JUN-2000; 2000WO-US016391

99US-0146194P

(WIST-) WISTAR INST ANATOMY & BIOLOGY

Halazonetis T, Scolnick D;

Novel nucleic acid sequence of mitotic checkpoint gene encoding a checkpoint with forkhead-associated domain and ring finger protein, diagnosing tumorigenic cells and in screening for anticancer drugs. for

ω --Page 38; 85pp; English

The present sequence is that of a 3' given in AAF30365, to amplify a cDNA nucleotides 1187-1753 of the human ch the human chfr sequence PCR primer, used with the 5' fragment corresponding to given primer The

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chfr gene encodes the human mitotic checkpoint protein Chfr (see AAB20219), which is required for regulation of the transition of cells from prophase to metaphase during mitosis. Loss of expression of Chfr is associated with a predisposition to tumourigenesis upon exposure to mitotic stress. A set of primers (see AAF30353-76) was used to amplify regions spanning the entire chfr coding region in order to determine whether the chfr gene is mutated in any of the human cancer cell lines $M8480, DLD1, HT29, HCT116, SAOS2, UZOS, IMR5 and NGP. A mutation leading to a Val-580 to Met amino acid substitution was identified in the chfr gene of UZOS cells. Chfr polypeptides and chfr nucleic acids are used in methods of diagnosing tumourigenic cells and of screening for drugs which can inhibit the activity of Chfr in a cancer cell, rendering it more
sensitive to additional antitumour therapies
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Sequence 18 BP; 6 A; 4 C; 5 G; 3 T; 0 U; 0 Other;

Query Match Best Local S Matches 18 l Similarity 18; Conserv Conservative 0; 3.6%; Score 18; Pred. No. Mismatches DB 1.2e+02; hes 0; 5 Length 18; Indels 0,

0

맑 Ś 179 18 CGTTTTGTGAGCTCAACC CGTTTTGTGAGCTCAACC 1 196

RESULT 5 AAF30370/ AAF30370 standard; DNA; 18

14-MAY-2001

Human checkpoint gene chfr 3'

(first entry)

PCR primer.

Checkpoint with forkhead associated domain and ring finger; Chfr; human; mitosis; cell cycle; tumour; diagnosis; antitumour; drug screening; ubiquitin-protein ligase; PCR primer; ss.

Homo sapiens

WO200109150-A2

14-JUN-2000; 2000WO-US016391

99US-0146194P

(WIST-) WISTAR INST ANATOMY & BIOLOGY

Halazonetis T, Scolnick D;

WPI; 2001-182927/18

Novel nucleic acid sequence of mitotic checkpoint gene encoding a checkpoint with forkhead-associated domain and ring finger protein, diagnosing tumorigenic cells and in screening for anticancer drugs.

Example 3; Page 38; 85pp; English

The present sequence is that of a 3' PCR primer, used with the 5' primer given in AAF30369, to amplify a cDNA fragment corresponding to nucleotides 1215-1753 of the human chfr sequence given in AAF30352. The chfr gene encodes the human mitotic checkpoint protein Chfr (see AAB20219), which is required for regulation of the transition of cells from prophase to metaphase during mitosis. Loss of expression of Chfr is associated with a predisposition to tumourigenesis upon exposure to mitotic stress. A set of primers (see AAF30353-76) was used to amplify regions spanning the entire chfr coding region in order to determine whether the chfr gene is mutated in any of the human cancer cell lines SW480, DLD1, HT29, HCT116, SAOS2, U2OS, IMR5 and NGP. A mutation leading to a Val-580 to Met amino acid substitution was identified in the chfr

Query Match Best Local Similarity

3.4%;

Score 17; Pred. No.

DB 5;

Length 17;

Sequence

17

B₽;

4 A;

7 C; 2

G; 4 T; 0 U; 0 Other;

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RESULT 6
AAF30362/c
ID AAF303
XX AAF303
XX AAF303
XX Checkp
XX Checkp
XX Checkp
XX Ubiqui
XX WO2001
XX WO2001
XX WO2001
XX WO2001
XX WIST-
PI 14-JUN
XX WFI;
PA (WIST-
XX WFI;
PA (WIST-
XX WFI;
PA (WIST-
XX WFI;
PA (WIST-
XX The pr checkp
PT checkp
PT checkp
PT checkp
PT checkp
PT checkp
PT checkp
CC Trom p
CC Chfr given
CC
                                              CC The present sequence is that of a 3' PCR primer, used with the 5' primer CC given in AAF30361, to amplify a cDNA fragment corresponding to CC nucleotides 904-1772 of the human chir sequence given in AAF30352. The CC chir gene encodes the human mitotic checkpoint protein Chir (see CL AAB20219), which is required for regulation of the transition of Chir is crown prophase to metaphase during mitosis. Loss of expression of Chir is associated with a predisposition to tumourigenesis upon exposure to CC mitotic stress. A set of primers (see AAF3033-76) was used to amplify cregions spanning the entire chir coding region in order to determine whether the chir gene is mutated in any of the human cancer cell lines CM substitution was identified in the chir code of U20S cells. Chir polypeptides and chir nucleic acids are used in methods of diagnosing tumourigenic cells and of screening for drugs which can inhibit the activity of Chir in a cancer cell, rendering it more can sensitive to additional antitumour therapies
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Halazonetis T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human checkpoint gene chfr 3' PCR primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14-JUN-2000; 2000WO-US016391
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-MAY-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sensitive to additional antitumour therapies
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         forkhead associated domaycle; tumour; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Scolnick D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BIOLOGY.
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antitumour;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chfr nucleic acids are used in and of screening for drugs which cer cell, rendering it more
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .2e+02;
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drug screening;
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human;

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RESULT 7
AAF30372/c
ID AAF303
XX AAF303
XX Human
XX Checkp
KW Mitosi
KW Ubiqui
XX Homo 8
XX WO2001
XX WIST-
XX WWI; 2
XX WWI; 
  RESULT 8
AAF30368/c
                                                                                                                                                                                                                                                                                                                                The present sequence is that of a 3' PCR primer, used with the 5' primer CC given in AAF30371, to amplify a cDNA fragment corresponding to nucleotides 1215-1772 of the human chfr sequence given in AAF30352. The Chir gene encodes the human mitotic checkpoint protein Chfr (see CC AAB20219), which is required for regulation of the transition of Cells from prophase to metaphase during mitosis. Loss of expression of Chfr is associated with a predisposition to tumourigenesis upon exposure to comitotic stress. A set of primers (see AAF3033-76) was used to amplify regions spanning the entire chfr coding region in order to determine whether the chfr gene is mutated in any of the human cancer cell lines SW480, DLD1, HT29, HCT116, SAOS2, UZOS, IMR5 and NGP. A mutation leading to a Val-580 to Met amino acid substitution was identified in the chfr gene of UZOS cells. Chfr polypeptides and chfr mucleic acids are used in methods of diagnosing tumourigenic cells and of screening for drugs which can inhibit the activity of Chfr in a cancer cell, rendering it more sensitive to additional antitumour therapies
                                                                                                                                                                                                                        Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel nucleic acid sequence of mitotic checkpoint gene encoding a checkpoint with forkhead-associated domain and ring finger protein, diagnosing tumorigenic cells and in screening for anticancer drugs.
                                                                                                                                                                                                                                                                                               Sequence 17
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                                                                                                                                                                                               . Similarity
17; Conserv
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                                                                                                                                                 GGTGACAAGTGTCTGGA 215
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                                                                                                                                                                                                                                                                                             BP; 4 A; 7 C; 2 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                     Score 17;
Pred. No.
                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                          Length 17;
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ADI25055
ID ADI2
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DT 22-A
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XX
KW domi
                                                                                                                                                                                                                                            RESULT 9
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                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Checkpoint with forkhead associated domain and ring finger; Chfr; h mitosis; cell cycle; tumour; diagnosis; antitumour; drug screening;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17
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                                                Human ABTB1
                                                                                              22-APR-2004
                                                                                                                                              ADI25055
                                                                                                                                                                                            ADI25055 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sensitive to additional antitumour therapies
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (WIST-) WISTAR INST ANATOMY & BIOLOGY
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17; Conserv
                                                                                                                                                                                                                                                                                                                                                               GGTGACAAGTGTCTGGA 215
                                             exon 3 reverse PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 4 A;
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                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                          3.3.5; F1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7 C; 2 G; 4 T;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 17;
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for

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dominant negative mutant RAB7; dominant negative mutant ARHGEF10;

Indels

<u>,</u>

Gaps

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ARESULT 10
AAF30375
ID AAF30
XX AAF30
XX AAF30
XX AAF30
XX Human
XX Human
XX Check
KW Check
KW mitos
KW ubiqu
XX Ubomo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CC The present invention describes an isolated nucleic acid (I) coding for a CC dominant negative, mutant RAB7 polypeptide and/or a dominant negative, mutant RAB7 polypeptide and/or a dominant negative, cc mutant ARHGEF10 polypeptide. (I) contains in comparison to the wild type RAB7 encoding sequence comprising 624 bp (SEQ ID NO: 1, AD125025) and/or CC to the wild type ARHGEF10 encoding sequence comprising 3366 bp (SEQ ID CC NO: 3, AD125027), one or more mutations, where the presence of the CC nucleic acids is indicative for a predisposition or presence of a CC peripheral neuropathy. Also described: (1) a nucleic acid probe which is CC a fragment of (I); (2) a recombinant vector comprising (I); (3) a host CC at fragment of a diagnostic assay to detect the presence of a peripheral neuropathy in a human; and (5) a transgenic non-human animal comprising CC the vector of (2). (I) is useful for isolating and detecting human CC peripheral neuropathy causing or predisposing genes. The diagnostic assay is useful for detecting the presence of peripheral neuropathy in a human. CC The present sequence represents a PCR primer for human ABFB1, which is
                                                                                                                                                                                                                                                                                                                                                                                                    Query Match

Best Local Similarity

Matches 16; Conserv
                                                                              Checkpoint with forkhead associated domain and ring finger; Chfr; human; mitosis; cell cycle; tumour; diagnosis; antitumour; drug screening; ubiquitin-protein ligase; PCR primer; ss.
                 WO200109150-A2
                                                Homo sapiens
                                                                                                                                                    Human
                                                                                                                                                                                   14-MAY-2001
                                                                                                                                                                                                                                                   AAF30375 standard; DNA; 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
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08-APR-2003;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            peripheral neuropathy; human; ABTB1; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (VLAA-) VLAAMS INTERUNIVERSITAIR INST BIOTECHNOG.
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                                                                                                                                                                                                                                                                                                                                                                      111
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                                                                                                                                                checkpoint gene
                                                                                                                                                                                                                                                                                                                                     ω
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    an example from the present invention.
                                                                                                                                                                                                                                                                                                                                                          CTGCCTGCAGCCTTTC 126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20
                                                                                                                                                                                                                                                                                                                                     CTGCCTGCAGCCTTTC 18
                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 2 A;
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2003EP-00076033.
                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           De
                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                  chfr
                                                                                                                                                                                                                                                                                                                                                                                                                                       3.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Jonghe
                                                                                                                                                  ບຸ
                                                                                                                                                                                                                                                                                                                                                                                                    Score 16; DB; Pred. No. 1.3
                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                              PCR primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Timmerman V,
                                                                                                                                                                                                                                                                                                                                                                                                                  DB 12;
. 1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Verhoeven
                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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RESULT 11
ABK69194/c
ID ABK691
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cc given in AAF30376, to amplify a cDNA fragment corresponding to comclectides 1625-2279 of the human chir sequence given in AAF30352. The chir gene encodes the human mitotic checkpoint protein Chfr (see CAAB20219), which is required for regulation of the transition of cells (from prophase to metaphase during mitosis. Loss of expression of Chfr is associated with a predisposition to tumourigenesis upon exposure to complicity regions spanning the entire chfr coding region in order to determine whether the chfr gene is mutated in any of the human cancer cell lines CC whether the chfr prince cid substitution was identified in the chfr gene of U2OS cells. Chfr polypeptides and chfr nucleic acids are used in the chfr gene of U2OS cells. Chfr polypeptides and chfr nucleic acids are used in methods of diagnosing tumourigenic cells and of screening for drugs which can inhibit the activity of Chfr in a cancer cell, rendering it more constitute to additional antitumour therapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 16; Conser
            modified_base
                                                                   modified_base
                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                         Antisense; phosphorylase kinase alpha 2; metabolic disorder; ss; antidiabetic; antiinflammatory; antimicrobial; cytostatic; diabetes;
                                                                                                                                                                                                                                                                                                                                    ABK69194;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel nucleic acid sequence of mitotic checkpoint gene encoding a checkpoint with forkhead-associated domain and ring finger protein, diagnosing tumorigenic cells and in screening for anticancer drugs.
                                                                                                                                                                                                                                                                                                      02-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                ABK69194
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 25 BP; 7 A; 9 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                  modified_base
                                                                                                                                                                                                             infection;
                                                                                                                                                                                                                                                                     Human phosphorylase kinase alpha 2 antisense oligonucleotide ISIS 118748
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence is that of a 5' PCR primer, used with the 3'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 3; Page 38; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Halazonetis T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-JUN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      118
                                                                                                                                                                                                                                                                                                                                                                                                                                          10
                                                                                                                                                                                                                                                                                                                                                                standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CAGCCTTTCTGCCACC
                                                                                                                                                                                                                                                                                                                                                                                                                                          CAGCCTTTCTGCCACC
                                                                                                                                                                                                             inflammation; tumour; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3.2%;
ilarity 100.0%;
Conservative (
                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                         /note=
1. .20
                          /note=
                                                                                                       _bom_
                                                                                                                                                 Location/Qualifiers
                                           /*tag= b
mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                DNA;
                                                                                       _base= OTHER
e= "Phosphorothioate backbone"
                             "All
                                                                                                                     D
a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     133
                                                                                                                                                                                                                                                                                                                                                                                                                                        25
                             cytosines
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                             are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.3e+03;
                             5-methylcytosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0,
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ADB6152/c
ID ADB615
XX ADB615
XX ADB615
XX ADB615
XX ADB615
XX Adrug,
KW C-Met
KW C-Met
KW Organı
KW Organı
KW W antiax
KW hepatiax
KW hepatiax
KW hepatiax
XX
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to an antisense compound 8-30 nucleobases in length targeted to a nucleic acid encoding phosphorylase kinase alpha-2, which specifically hybridizes with and inhibits the expression of phosphorylase kinase alpha-2. The antisense compound is useful for inhibiting the expression of phosphorylase kinase alpha-2 in human cells or tissues, and for treating a human having a disease or condition associated with phosphorylase kinase alpha-2 including metabolic disorder or diabetes. The antisense compound is useful for diagnostics, therapeutics, prophylactically to prevent or delay infection, inflammation or tumour formation. The antisense compound is safely and effectively administered to humans. The present sequence is an antisense oliponucleotide of the
                                            c-Met regulatory mucin; Mermuc; HGF-dependent cell proliferation; organ; blood vessel; mucous membrane; bone; nerve formation; regeneration; neogenesis; hepatotropic; nephrotropic; antiarteriosclerotic; antiinflammatory; virucide; liver; hepatic sclerosis; hepatic fibrosis; symptomatic; acute; virikidney; chronic kidney disease; lung; arteriosclerosis; PCR;
                                                                                                           drug; binding inhibitor; hepatocyte growth factor recptor; HGF; c-Met; c-Met regulatory mucin; Mermuc; HGF-dependent cell proliferation; tiss organ; blood vessel; mucous membrane; bone; nerve formation; repair;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 4 A; 7 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense compounds targeted to nucleic acids encoding phosphorylase kinase alpha-2, useful for modulating gene expression and treating diseases associated with expression of phosphorylase kinase alpha-2 in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-SEP-2000; 2000US-00657453
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-MAR-2002
                                                                                                                                                                                                                                                                      ADB61552
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                                                                                                                                                                      Hepatocyte growth factor (HGF)
                                                                                                                                                                                                         04-DEC-2003
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                  unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Page
                                                                                                                                                                                                                                                                      standard;
                                                                                                                                                                                                                                                                                                                                                                                 CCTGTGCACCTGCTG
                                                                                                                                                                                                                                                                                                                                                    CCTGTGCACCTGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             targetting human phopsphorylase kinase alpha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Wyatt JR;
                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PHARM INC
                                                                                                                                                                                                         (first entry)
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16. .20
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e= "2'-O_methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "2'-0_methoxyethyl residues"
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                                                                                                                                                                                                                                                                                                                                                                                                                                Score 15;
Pred. No.
                                                                                                                                                                        receptor related primer,
                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                 No.
                                                                                                                                                                                                                                                                                                                                                                                                                                DB 6; Lo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 20
                                                                                                                                                                          SEQ
                                              PCR; primer;
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                                                                                                                                                                                                                                                                                                                                                                                                                 0,
                                                                                                         repair;
                                               hepatitis;
rimer; ss.
                                                                                                                                                                          No 25
                                                                                                                             tissue;
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Best Local
                                                                                                                     gene therapy; Lambert-Eaton myasthenic syndrome; LEMS; PCR; autoimmune disease; epilepsy; migraine; episodic ataxia; cancer; strr brain trauma; Alzheimer's disease; multiinfarct dementia; convulsion; Korsa-koff's disease; amyotrophic lateral sclerosis; seizure; Huntington's disease; amnesia; cardiac arrhythmia; angina pectoris; hypoxia; ischaemia; myocardial infarction; congestive heart failure; muscular dystrophy; hypertension; chromosome 3p21.3; lung cancer; breacplastic lesion; hyperplasia; dysplasia; carcinom ascet cancer; preneoplastic lesion; hyperplasia; dysplasia; carcinom
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a novel drug composition containing a substance active in inhibiting the binding of the hepatocyte growth factor (HGF) receptor c-Met to c-Met regulatory mucin (Mermuc) and in doing so regulates HGF-dependent cell proliferation and tissue, organ, blood vessel, mucous membrane, bone and nerve formation, repair, regeneration and neogenesis. The novel drug composition has the following activities: hepatotropic, nephrotropic, antiarteriosclerotic, antiinflammatory, and virucide. The drug composition is useful for the treatment and prevention of diseases of the liver (including hepatic sclerosis, hepatic fibrosis, and symptomatic, acute or viral hepatitis), kidney (including chronic kidney disease) and lung, and arteriosclerosis. This polynucleotide sequence represents the DNA encoding an HGF receptor of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Drug composition containing regulator of binding of HGF receptor a regulatory mucin for treatment and prevention of liver, kidney disease by regulation of cell proliferation and tissue formation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13-DEC-2001; 2001JP-00380158
04-DEC-2002; 2002JP-00352924
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                                                                                                                                                                                                                                                                                             Human calcium channel alpha2delta SSCP PCR primer #24.
                                                                                                                                                                                                                                                                                                                               09-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                ABS74300
                                                                                                                                                                                                                                                                                                                                                                                               ABS74300 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12-DEC-2002; 2002WO-JP013014
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CAGCCTTTCTGCCAC 132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ζ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CAGCCTTTCTGCCAC
                                                                                                                                                                                                                                                          primer; calcium channel alpha2delta; splice isoform; CACNA2D2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 7 A; 3 C; 8 G; 3 T; 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Page 198; 223pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Higuchi
                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                            polymorphism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 15;
Pred. No
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4.1e+03;
hes 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                            convulsion;
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27-AUG-2002. US6441156-B1

sapiens

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ACD25698
ID ACD2
XX
AC ACD2
XX
AC ACD2
XX
XX
C BC FUMBE
XX
DE Huma
XX
KW Huma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a purified nucleic acid sequence (referred as CACNA2D2 gene which encodes human calcium channel alpha2delta-2 subunit CC protein) comprising a fully defined alpha2delta splice isoform 1, 2 or 3 cc nucleic acid sequence, or its complement and the encoded protein which C include are: (1) a method of producing a calcium channel protein which C involves introducing a recombinant expression vector comprising the CC CACNA2D2 nucleic acids and encoding the calcium channel protein, into a cultured host cell under conditions such that the host cell expresses the CC amino acid sequences; and (2) a method for co-expressing calcium channel proteins, comprising carrying out the method of (1), but with one or more cc producing a calcium channel protein acid sequences and comprising one or more nucleic acid sequences cc producing the splice variants. CACNA2D2 nucleic acid is useful for protein activity (e.g. epilepsy migraine, episodic ataxia, cancer, stroke, brain trauma, Alzheimer's disease, multiinfarct dementia, cancer, stroke, brain trauma, angenesia, cardiac arritythmia, angina cancer, stroke, brain trauma, angenesia, cardiac arritythmia, angina cancer, stroke, st
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         preneoplastic lesion (e.g. hyperplasia, dysplasia, carcinoma in situ). The present is an SSCP (single strand change polymorphism) PCR primer used to detect polymorphisms in sequences encoding a human calcium channel alpha2delta splice isoform protein
                                                             Human calcium channel alpha2delta SSCP primer MJE7R.
                                                                                                                              26-AUG-2003
                                                                                                                                                                                                                                                         ACD25698 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel purified nucleic acid sequence encoding human calcium channel alpha2delta subunit protein, useful for detecting, preventing and treating cancer, stroke, brain trauma, Huntington's disease, myocar
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PCR; calcium channel alpha2delta; chromosome 3p21.3; primer;
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CC subunit splice isoform 1, splice isoform 2 sequence or splice isoform 3, or their variants, and their encoding nucleic acids (or their variants, and their encoding nucleic acids (or their variants, and their encoding nucleic acids (or their compound for modulating calcium channel activity, an antibody which binds compound for modulating calcium channel activity, an antibody which binds chuman animal (where the animal expresses a reduced level of calcium channel alpha 2delta subunit relative to a corresponding wild-type canimal. The calcium channel proteins are useful for generating an cantibody (which is useful for detecting the proteins or their portions). The transgenic animal (preferably a rodent e.g. mouse) is useful for calcium channel proteins are useful for generating an cantibody (which is useful for detecting the proteins or their portions). The carting a therapeutic compound for treating a transgenic animal cantibody (which is useful for detecting the proteins or their portions). The having cancer, especially lung cancer (small call carcinoma or non-small cancer, especially carciant (small call carcinoma or non-small cancer, cell carcinoma or non-small cancer, campotrophic lateral sclerosis, convulsions, seizures, Huntington's tracke, brain trauma, Alaheimer's disease, multiinfarct dementia, convulsions, seizures, Huntington's convulsions, seizures, Huntington's carchythmia, angina pectoris, hypoxic damage to the cardiovascular system, and congestive heart failure, or Lambert-Eaton mysachenic syndrome. The calcium channel alpha2delta subunit gene for the calcium channel alpha2delta subunit gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New substantially pure human calcium channel alpha2delta subunit splice isoform 1, 2 and 3 sequence useful in preventing, treating and diagnosing cancer, neurological disorders and cardiovascular disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          amyotrophic lateral sclerosis; convulsions; Huntington's disease; amnesia; cardiovascular disease; cardiac arrhythmia; angina pectoris; hypoxic damage; ischaemia; myocardial infarction; SSCP; congestive heart failure; Lambert-Baton myasthenic syndrome; single strand conformation polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        transgenic; cancer; lung cancer; small cell carcinoma; epilepsy; s
non-small cell carcinoma; breast cancer; nasopharyngeal cancer;
cervical cancer; head and neck cancer; neurological disease;
brain trauma; Alzheimer's disease; multiinfarct dementia; seizure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         comprising at least
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a substantially purified amino acid
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WEI M.
DUH F.
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SEKIDO Y.
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99US-00470443.
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                                                                                                                                                                                                                                                                                                           The sequence represents the Afx transcription factor response element, selected sequence #12. Human fork head transcription factor, Afx, was expressed and the protein used to select a response element comprising an base pair (bp) nucleotide sequence. The nucleotide sequence comprises AACATGTT, the Afx response element, and is useful in bioinformatic analysis e.g. of the human genome. Employing the Afx response element is also useful for screening for genes that may be used as diabetes drug targets. This can provide a subset of genes transcriptionally responsive to insulin and may lead to development of assays that facilitate the analysis of genes interacting with the insulin receptor pathway. The genes found in such screening may also be employed in further screening methods for compounds which modify the insulin receptor signalling
                                                                                                                                                                                                                                   Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New Afx response element with a nucleotide sequence comprising a DNA binding site for the human fork head transcription factor Afx, useful screening for genes or in bioinformatic analysis of the human genome.
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31-AUG-1999;
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RESULT 16 & ACI20807 Standard; DNA; 25
                                                                           The invention discloses a microarray comprising a plurality of nucleic CC acid probes including one of 2,018,500 fully defined sequences, or its comperfect match, perfect mismatch, antisense match or antisense mismatch. CC Also disclosed is a method of gene expression analysis. The array is used CC in monitoring gene expression levels by hybridisation to a DNA library, CC in analysis of genetic variation or in hybridisation to a DNA library, CC in analysis of genetic variation or in hybridisation to a DNA library, CC in analysis one target sequence. The method of analysis comprises CC hybridising at least one or more nucleic acids to at least two or more cC nucleic acid probes and detecting the hybridisation. The nucleic acid grobes are attached to a solid support. The analysis comprises monitoring CC gene expression levels, identifying biallelic markers or polymorphisms, CC or family members of a gene and a cross-species comparison. Each of the nucleic acids further comprises a tag sequence. The array of nucleic acid probes is useful in in situ hybridisation, in Southern, Northern or dotolot hybridisation to identify or detect the sequence or specific cutations of any gene, in mapping the 5' termini of mRNA molecules by primer extensions or in screening cDNA or genomic libraries or subclones CC for additional subclones containing segments of DNA that have been contained and previously sequenced. The sequence presented is one of the nucleic acid probes incorporated in the microarray. Note: The sequence CC data for this patent can also be obtained in electronic format directly cx
                   Query Match
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                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New array of nucleic acid probes, useful for in Southern, Northern or dot-blot hybridization to sequence or specific mutations of any gene.
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genetic variation;
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on; biallelic marker; polymorphism; human;
               100.0%;
                                                                           5 C; 6
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                   Score 15; DB 9;
Pred. No. 4.1e+0
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                   4.1e+03;
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                                     Length 25;
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ACH57351

ACH57351

ACH57351

ACH57251

ACH67251

ACH72251

ACH722
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                                                                                                                                                                                                                                                                                                                      CC complementary to particular genes, and can be used as probes for a CC complementary to particular genes, and can be used as probes for a CC variety of analyses such as gene expression analysis. Each probe CC comprises 9 or more consecutive nucleotides from at least one of 14936 CC nucleotide sequences defined in the patent, or their perfect sense match, antisense match or antisense mismatch oligonucleotides. The probes may be used in an array comprising at least 10 distinct CC nucleic acid probes. The array is useful in monitoring gene expression CC levels by hybridisation to a DNA library, in analysing genetic variations, and in hybridising tag-labelled compounds. The probes are useful for identifying family members of a gene. The probes are also cuseful in in situ hybridisations, in screening cDNA or genomic libraries (or derived subclones) for additional clones containing segments of DNA that have been previously isolated and sequenced, in Southern, northern, CC that have been previously isolated and sequenced, in Southern, northern, CC mapping the 5' termini of mRNA molecules by primer extensions. The nucleic acid sequences of the invention are also useful as PCR primers. CC complementary to particular genes with a wide range of analytical uses. CC complementary to particular genes with a wide range of analytical uses. CC The sequence data for this patent was obtained in electronic format cromat directly from the USPTO web site at segdata.uspto.gov/psipsDIDEntry.html
                                                                                                                                                                  Query Match
Best Local
                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New probe array useful e.g. for monitoring gene expression levels, analyzing genetic variations, or for hybridizing tag-labeled compoundables multiple nucleic acid probes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08-AUG-2001; 2001US-0311040P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                tag-labelled compound; gene family; in situ hybridisation; library screening; Southern hybridisation; northern hybridisation; dot-blot hybridisation; gene sequence; mutation detection; target sequence; probe; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gene expression analysis; array; hybridisation; genetic variation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA target sequence #6487 useful in array for genetic analyses
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-OCT-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ACH57351 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-576608/54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mittmann
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; SEQ ID NO 6487; 9pp; English.
                                                                   156
                                                                                                                                      15;
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                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                        25
                                                               CGGCTGCTACGGCTG 170
CGGCTGCTACGGCTG 22
                                                                                                                                                                                                                                                                     BP; 6 A; 6 C; 9 G; 4 T; 0 U;
                                                                                                                                      Conservative
                                                                                                                                                                  3.0%;
100.0%;
                                                                                                                                  0,
                                                                                                                                                                      Score 15;
Pred. No.
                                                                                                                                      Mismatches
                                                                                                                                                                  DB 9; L
                                                                                                                                                                                                                                                                     0 Other;
                                                                                                                                                                                                  Length 25
                                                                                                                               Indels
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RESULT 18

Query Match
Best Local Similarity 78.0
Matches 11; Conservative

2.8%;

Score 14; DB 2; Length 15; Pred. No. 1.3e+04; 3; Mismatches 0; Indels

0

Gaps

0

15

BP; 5 A;

3 C; 4 G; 0 T;

3 U; 0 Other;

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The present invention describes a novel enzymatic nucleic acid (ENA) CC having a hammerhead motif (HM) comprising; (i) at least 5 ribose residues (; (ii) a 2'-C-allyl modification at position 4 of the ENA; (iii) at least 5 cc can inhibit collagenase and stromelysin production in the synovial CC membrane of joints for the treatment or prevention of arthritis, can also be used to treat antigen presenting cells of a donor to induce tolerance in a recipient to an alloantigen of a donor. They can also be used for creating graft tolerance or for treating autoimmune disease, and for treating allergies and other inflammatory conditions. The ENA's can also be used in diagnosis. Ribozyme therapy impacts on the expression of cromelysin without introducing the non-specific effects upon gene expression which accompany treatment with retinoids and dexamethasone. The present sequence is used in the exemplification of the present sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-APR-1995;
02-MAY-1995;
04-MAY-1995;
07-JUL-1995;
07-JUL-1995;
07-AUG-1995;
05-OCT-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-DEC-1994;
23-DEC-1994;
23-DEC-1994;
17-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Beigelman
Mcswiggen
Karpeisky
                                                                                                                                                                                                                                                                                                                                                   Enzymatic nucleic acid molecules having a hammer-head motif - used for the treatment of arthritis, induction of graft tolerance or treatment of auto-immune diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Arthritic condition; graft tolerance; immune response; target; cleavage; hammerhead ribozyme; hairpin ribozyme; huanna; rabbit; mouse; collagenase; stromelysin; synovial membrane; joint; arthritis; ostcoarthritis; rheumatoid arthritis; autoimmune disease; allergy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mus sp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAX65296;
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                               present invention
                                                                                                                                                                                                                                                                                                                        Claim 10;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                       Page 179; 307pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                          Stinchcomb DT, Jarvis T, Draper K, Gustofson J, Usman N, Wincott F, I Thompson JD, Modak A, Burgin A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     94US-00354920
94US-00363253
94US-00363254
95US-00390850
95US-00432874
95US-00432874
95US-00434509
95US-0000971P
95US-0000971P
95US-0000971P
95US-0000971P
95US-0000974
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RESULT 19
AAX65298
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The present invention describes a novel enzymatic nucleic acid (ENA) Chaving a hammerhead motif (HM) comprising: (i) at least 5 ribose residues (; (ii) a 2'-C-allyl modification at position 4 of the ENA; (iii) at least ten 2'-O-methyl modifications; and (iv) a 3'-end modification. The ENA's C can inhibit collagenase and stromelysin production in the synovial membrane of joints for the treatment or prevention of arthritis, can also be used to treat antigen presenting calls of a donor to induce tolerance in a recipient to an alloantigen presenting autoimmune disease, and for C treating altergies and other inflammatory conditions. The ENA's can also be used in diagnosis. Ribozyme therapy impacts on the expression of C stromelysin without introducing the non-specific effects upon gene cappression which accompany treatment with retinoids and dexamethasone. The concentration of ribozyme required to affect a therapeutic treatment C I lower than that required of antisense molecules, and is highly considered. The present sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-DEC-1994
23-DEC-1994
23-DEC-1994
17-FEB-1995
20-APR-1995
02-WAY-1995
04-WAY-1995
07-JUL-1995
07-JUL-1995
                                                                                                                                                                                                                                                                                                                                                           Beigelman
Mcswiggen
Karpeisky
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Arthritic condition; graft tolerance; immune response; target; cleavage; hammerhead ribozyme; hairpin ribozyme; human; rabbit; mouse; collagenase; stromelysin; synovial membrane; joint; arthritis; osteoarthritis; rheumatoid arthritis; autoimmune disease; allergy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       diagnosis;
                                                                                                                                                                                                                                                                                         Enzymatic nucleic acid molecules having a hammer-head motif
the treatment of arthritis, induction of graft tolerance or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9618736-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mouse
                                                                                                                                                                                                                                                     Claim
                                                                                                                                                                                                                                                                               auto-immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22-NOV-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                   (RIBO-)
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                                                                                                                                                                                                                                                    10; Page 179; 307pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N
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                                                                                                                                                                                                                                                                               diseases
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Gustofson J, U
Thompson JD, M
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                                                                                                                                                                                                                                                                                                                                                                                                                                            95US-00390850
95US-00426124
95US-00432874
95US-00434509
95US-0000974P
95US-0000974P
95US-00512861
95US-00541365.
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94US-00363254.
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                                                                                                                                                                                                                                                                                                                                                                                                                    PHARM INC
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                                                                                                                                                                                                                                                                                                                                                             Usman
Modak
                                                                                                                                                                                                                                                                                                                                                           Jarvis T, Draper K,
sman N, Wincott F, I
odak A, Burgin A;
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                                                                                                                                                                                                                                                                                                                                                                           Pavco P;
Matulic-Adamic

    used for
treatment

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the

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RESULT 20
AAX65297
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Best Local S
Matches 11
                                                                                                                                                                                                                                                                               20-APR-1995;
02-MAY-1995;
04-MAY-1995;
07-JUL-1995;
07-JUL-1995;
07-AUG-1995;
05-OCT-1995;
                                                                                                                                                                                                                                                                                                                                                              13-DEC-1994;
23-DEC-1994;
23-DEC-1994;
17-FEB-1995;
          The present invention describes a novel enzymatic nucleic acid (ENA) having a hammerhead motif (HM) comprising: (i) at least 5 ribose residues; (ii) a 2'-C-allyl modification at position 4 of the ENA; (iii) at least ten 2'-C-methyl modifications; and (iv) a 3'-end modification. The ENA's can inhibit collagenase and stromelysin production in the synovial membrane of joints for the treatment or prevention of arthritis, particularly osteoarthritis or rheumatoid arthritis. The ENA's can also be used to treat antigen presenting cells of a donor to induce tolerance in a recipient to an alloantigen of a donor. They can also be used for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 15 BP; 5
                                                                                                                                                                                                              Mcswiggen
Karpeisky
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Arthritic condition; graft tolerance; immune response; target; oll hammerhead ribozyme; hairpin ribozyme; human; rabbit; mouse; coll stromelysin; synovial membrane; joint; arthritis; osteoarthritis; rheumatoid arthritis; autoimmune disease; allergy; inflammation;
                                                                                                                                                        Enzymatic nucleic acid molecules having a hammer-head motif - used the treatment of arthritis, induction of graft tolerance or treatme
                                                                                                                                                                                                                                                                                                                                                                                                                                            20-JUN-1996.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           diagnosis;
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                                                                                                                         Claim 10;
                                                                                                                                                                                                                                     Beigelman
                                                                                                                                                                                                                                                           (RIBO-)
                                                                                                                                                                                                                                                                                                                                                                                                                      22-NOV-1995;
                                                                                                                                             treatment of arth
o-immune diseases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard;
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graft tolerance or
                                                                                                                      Page 179; 307pp;
                                                                                                                                                                                                               ÞGF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             hammerhead ribozyme target SEQ ID
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94US-00363254.

95US-00390850.

95US-00426124.

95US-00432874.

95US-00009517.

95US-0000974P.

95US-00512861.

95US-00541365.
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                                                                                                                                                                                                                                                           PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.8%;
78.6%;
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הֻ מָּ
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Modak
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Pred. No. 1
 for treating autoimmune
                                                                                                                                                                                                              Jarvis
Jsman N,
Modak A,
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Burgin A;
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                                                                                                                                                                                                                           Matulic-Adamic
                                                                                                                                                                                                                                       Pavco
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                                                                                                                                                          treatment
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RESULT 21
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Best Local
The present invention relates to nucleic acid probes, which are useful for assaying nucleic acids by hybridising with a target nucleic acid, in which a single-stranded oligonucleotide is labelled with a fluorescent substance and a quencher in a manner that the fluorescence intensity of the hybridisation reaction system is increased after completion of the hybridisation but no stem loop structure is formed. The probes are useful for assaying nucleic acids and their polymorphism and mutation, particularly useful for e.g. analytical applications, disease diagnosis and microbial identification. The present sequence was used to illustrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27-JUN-2000;
03-AUG-2000;
26-SEP-2000;
                                                                                                                                                                                                                                                                                    Fluorescently-labeled nucleic acid probes for assaying nucleic acids their polymorphism and mutation, particularly useful in science and medicine for e.g. analytical applications, disease diagnosis and microbial identification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  treating allergies and other inflammatory conditions. The ENA's can also be used in diagnosis. Ribozyme therapy impacts on the expression of stromelysin without introducing the non-specific effects upon gene expression which accompany treatment with retinoids and dexamethasone. The concentration of ribozyme required to affect a therapeutic treatment is lower than that required of antisense molecules, and is highly specific. The present sequence is used in the exemplification of the
                                                                                                                                                                                                                                             Example 41;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kurane R,
Yokomaku T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27-JUN-2001;
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                                                                                                                                                                                                                                      Page 103; 152pp;
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Matches 14
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Best Local (
                                                                                                                            invention are also useful in the treatment of: inflammatory bowel disease, glutenenteropathy, infectious diseases, autoimmune diseases (e.g. haemolytic anaemia, rheumatoid arthritis, dermatitis, Grave's disease, multiple sclerosis, allergy, asthma and diabetic mellitus), diseases or disorders of the immune system, hypersensitivity, anaphylaxis, and blood group incompatibility. The present DNA sequenc represents a PCR primer that was used to amplify an intestinal epithelium/peyer's patch M cell-associated DNA sequence of the invent
                                                                                                                                                                                                                                         The invention comprises DNA sequences which eals are associated with intestinal epithelium and peyer's patch M cells. The DNA sequences of the invention are useful for assessing, modifying, modulating or regulating intestinal epithelium or M cell development. The DNA sequences of the
                                                                                                                                                                                                                                                                                                                                 Novel isolated or purified polypeptide encoded by genes associated with intestinal epithelium or M cell development, differentiation or function useful for treating autoimmune diseases and infectious diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         inflammatory bowel disease; glutenenteropathy; infectious disease; autoimmune disease; haemolytic anaemia; rheumatid arthritis; dermatitis; Grave's disease; multiple sclerosis; allergy; asthma; diabetic mellitus; immune system disorder; hypersensitivity; anaphylaxis;
                                                                                                                                                                                                                                                                                                          Disclosure; SEQ ID NO 539; 152pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                      Brayden DJ,
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lity; ss; human; PCR; primer.
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                                                               1.3e+04;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to antisense compounds, especially oligonuclectides, which are targeted to a nucleic acid encoding inducible nitric oxide synthase and which specifically hybridise to and modulate expression of inducible nitric oxide synthase. The antisense compounds have immunomodulator, antidiabetic, cardiovascular, cardiant, neuroprotective, disorder and vasotropic activity. The antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        oligonucleotides are useful for inhibiting the expression of inducible nitric oxide synthase in cells or tissues. In particular, the antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense oligonuclectides for modulating the expression of inducible nitric oxide synthase in cells or tissues, particularly useful for treating e.g. immunological, cardiovascular or neurological disorders, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     modulate expression; immunomodulator; antidiabetic; cardiovascular; cardiant; neuroprotective; vasotropic; ischaemia; reperfusion injury;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 residues are 5-methylcytidines and a deoxy gap"
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                                                                                                                                                                                                                                                                      gequences of the invention contain numerous binding sites for members of the HMG-I(Y) ("high mobility group") family of nonhistone chromatin corrections, a group of minor groove-binding architectural transcription factors which are thought to be involved in the mechanisms by which EASE sequences improve expression of transgenes. The EASEs of the invention can also be used in the identification of additional EASE sequences (e.g. from other, transformed cell lines which exhibit high levels of expression not attributable to a high gene copy number). Expression of recombinant therapeutic proteins in mammalian cells is often preferable to expression in microbial (prokaryotic) cells, since the post-translational modifications found in mammalian cells are more likely to resemble those found in a mammal. Sequences AAX98348 - AAX98353 represent PCR primers used to create electrophoretic mobility shift assay (EMSA) probes
                                                                                                              Matches
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13-JAN-1997;
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                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention comprises Chinese hamster expression augmenting sequence elements (EASEs; AAK98343-AAK98344) that can be used to improve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-OCT-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-033281/04.
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                                                       TGTTTCTGCTGTCT 345
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                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Es; AAK98343-AAK98344) that can be used to improve recombinant proteins in mammalian host cells. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                96US-00586509.
97US-00785150.
99US-00435377.
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                               ADM36243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to an isolated polynucleotide comprising a nucleic acid molecule comprising nucleotides 11538-11692, nucleotides 11538-11760, nucleotides 11673-12165, nucleotides 11813-12165 or nucleotides 11899-12165 of ADD89798, the hamster high mobility group, HMG-I(Y) gene, fragments of the DNA having expression augmenting activity (an expression augmenting sequence element, EASE) or their combinations or complementary DNA. Also included are a mammalian host cell which comprises the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           expression augmenting sequence element; EASE; EMSA; electrophoretic shift mobility assay.
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05-NOV-1999; 99US-00435377.
02-MAR-2000; 2000US-0186537P.
12-SEP-2000; 2000US-00660299.
                                                                                ADM36243
                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
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Best Local S
Matches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention relates to novel membrane proteins that are associated with differentiation and/ or metabolic function of adipocytes, in particular of mouse origin. Specifically, it refers to the isolated nucleic acid molecules encoding all or part of these proteins, appropriate antibodies and screening assays useful for the development of drug compositions derived thereof. The present invention describes these compositions as useful for the treatment of diseases associated with abnormalities of adipocyte function, such that they can be used to prevent, treat or diagnose obesity, hypertension, hyperlipaemia, diabetes and arteriosclerosis. Accordingly, they exhibit anorectic, antidiabetic, antiarteriosclerosic, antilipaemic and hypotensive activities. This oligonucleotide is a 3' RACE gene specific PCR primer used to amplify murine adipocyte-derived cDNA of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCR; 88; obesity;
                                              antiarteriosclerotic; neuroprotective; nootropic; antiparkinsonian; anticonvulsant; pulmonary disease; restenosis; atherosclerosis; Alzheimer's; Parkinson's; epilepsy; dementia; huntington's;
Unidentified
                                amyotrophic
                                                                                               double-stranded short interfering nucleic acid; siNA;
                                                                                                                                  EGFR-targeted sinA DNA-RNA hybrid -
                                                                                                                                                                   26-FEB-2004
                                                                                                                                                                                                                                     ADG29799 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mouse membrane and secretory proteins of adipocyte origin and polynucleotides encoding them for screening compounds as remedies for obesity, diabetes, arteriosclerosis, hypertension and hyperlipemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3' RACE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 3; SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-JUL-2003; 2003WO-JP008690
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           03-JUN-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matsuzawa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-JUL-2002; 2002JP-00201856
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22-JAN-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO2004007711-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          anorectic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (TAKE )
                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2004-122943/12.
                                                                                                                                                                                                                                                                                                                                                    28
                                                                                                                                                                                                                                                                                                                      N
                                                                                                                                                                                                                                                                                                                                                                                     14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TAKEDA CHEM IND
                                                                                                                                                                                                                                                                                                                                                                                                                                                       20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        mouse; murine; differentiation; metabolic function; adipocyte;
hypertension; hyperlipaemia; diabetes; arteriosclerosis;
c; antidiabetic; antiarteriosclerotic; antilipaemic; hypotensive;
                                                                                                                                                                                                                                                                                                                                                    TGCACCTGCTGCTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ķ
                                lateral
                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 3 A; 7 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          primer used to
                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Funahashi T, Shimomura C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ID NO 38; 195pp; Japanese.
                                                                                                                                                                                                                                   RNA;
                               sclerosis; gene
                                                                                                                                                                                                                                                                                                                                                                                                                        2.8%;
                                                                                                                                                                                                                                     21
                                                                                                                                                                                                                                                                                                                                                    41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LTD.
                                                                                                                                                                                                                                     ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          amplify murine SST8-5
                                                                                                                                                                                                                                                                                                                                                                                     <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                   Score 14;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                               therapy; ss; DNA-RNA hybrid; EGFR
                                                                                                                                  SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                      DB 12;
                                                                                                                                                                                                                                                                                                                                                                                                     1.3e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Furuyama
                                                                                                                                                                                                                                                                                                                                                                                                                  Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                     Indels
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RESULT 28
ADL80074/c
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                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20-FEB-2002;
11-MAR-2002;
06-JUN-2002;
29-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                   useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary disease such as restenosis, atherosclerosis, Alzheimer's disease, Parkinson's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Furthermore, the siNA may be utilised for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a double-stranded short interfering nucleic (siNA) molecule that down-regulates expression of an endogenous mamma target gene comprising one or more chemical modifications and each st of the double-stranded siNA comprises about 21 nucleotides. The siNA the invention demonstrates antiarteriosclerotic, neuroprotective, neuroprotective, antiparkingonian and anticonvalsant activities and may be
RNA interference; short interfering nucleic acid; siNA; short interfering RNA, siRNA, double-stranded RNA, micro-RNA, miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cancer; cytostatic; human; oncogene; epidermal growth factor receptor; EGFR;
                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene of treating diseases that respond to modulation of gene expression or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-731676/69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-SEP-2002;
09-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-FEB-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO2003074654-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example
                                                                                                  Human HER1
                                                                                                                                  20-MAY-2004
                                                                                                                                                                                        ADL80074
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (SIRN-) SIRNA THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                        therapy applications. The current sequence is
                                                                                                                                                                                                                                                                              244 TCAGACATCCTGAA 257
                                                                                                                                                                                                                                                                16
                                                                                                                                                                                                                                                                                                                       14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24; SEQ ID NO 365;
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                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                              21
                                                                                                                                                                                        standard;
                                                                                                                                                                                                                                                                TCAGACATCCTGAA 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ç
                                                                                                                                                                                                                                                                                                                                                                                                         of the invention.
                                                                                                    (EGFR) chemically modified siRNA, SEQ ID NO:1241.
                                                                                                                                                                                                                                                                                                                                                                             BP; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2002US-0358580P.
2002US-0363124P.
2002US-0386782P.
2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
                                                                                                                                                                                                                                                                                                                    2.8%; So
larity 100.0%; I
Conservative 0;
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                                                                                                                                (first entry)
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                                                                                                                                                                                                                                                                                                                                                                             RNA;
                                                                                                                                                                                                                                                                                                                                                                             2 C; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Thompson J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ۲
                                                                                                                                                                                         21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        593pp; English.
                                                                                                                                                                                                                                                                                                                     Score 14; DB; Pred. No. 1.3
                                                                                                                                                                                                                                                                                                                                                                             G; 2 T; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chowrira B,
                                                                                                                                                                                                                                                                                                                                                                             U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                   DB 10;
                                                                                                                                                                                                                                                                                                                                   1.3e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pavco
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D
                                                                                                                                                                                                                                                                                                                                               Length 21;
                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                          that of
                                                                                                                                                                                                                                                                                                                                                                                                                          the siNA DNA-
                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    each strand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cleic acid
                                                                                                                                                                                                                                                                                                                    Gaps
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20-FEB-2002; 2002US-035B580P.
11-MAR-2002; 2002US-033124P.
29-MAY-2002; 2002US-016840.
06-JUN-2002; 2002US-0113552.
06-JUN-2002; 2002US-0113552.
06-JUL-2002; 2002US-0393924P.
29-AUG-2002; 2002US-0406784P.
29-AUG-2002; 2002US-0408378P.
05-SEP-2002; 2002US-0408378P.
19-SEP-2002; 2002US-040939P.
19-SEP-2002; 2002US-040293P.
19-SEP-2002; 2002US-00277494.
15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                20-FEB-2003; 2003WO-US005045
                                                                                                                                                                                                                                                                                                                                                                       modified_base
                                                                                                                                                                                                                                                                                                                                                                                                   modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                               modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HER1; c-erb-B-1; DNA-RNA hybrid; phosphorothioate; ss
                                                                                                                                                                                                                                                                                                               28-AUG-2003
                                                                                                                                                                                                                                                                                                                            WO2003070912-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                              modified_base
                                                                                                                                                                                                                                                                                                                                                                 /*tag= d
/mod_base= OTHER
/mote= "2'-deoxy-2'-fluoro base"
                                                                                                                                                                                                                                                                                                                                                                                                                                             /*tag= b
/mod_base=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
3...5
                                                                                                                                                                                                                                                                                                                                                                                                                   /*tag= c
/mod_base=
                                                                                                                                                                                                                                                                                                                                                                                                     /note=
19
                                                                                                                                                                                                                                                                                                                                                /*tag= e
/mod_base= OTHER
/note= "Ribothymidine.
                                                                                                                                                                                                                                                                                                                                                                                                                                        note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mod
                                                                                                                                                                                                                                                                                                                                          phosphorothicate"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            base=
                                                                                                                                                                                                                                                                                                                                                                                                            "2'-deoxy-2'-fluoro bases'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "2'-deoxy-2'-fluoro bases"
                                                                                                                                                                                                                                                                                                                                                                                                                                         -deoxy-2'
                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                         -fluoro
                                                                                                                                                                                                                                                                                                                                                  Also, the internucleotide linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                         base"
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(RIBO-) RIBOZYME PHARM INC.

Mcswiggen 4 Pavco P, Beigelman L, Fosnaugh K, Jamison

s

WPI; 2003-697612/66

New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of the epidermal gro receptor gene epidermal growth

Example 9 SEQ ID NO 1241; 171pp; English

downregulate expression of one or more human epidermal growth factor receptor (EGFR) genes (including HER1, HER2 HER3 and HER4) by RNA interference. The siNAs may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA) double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are The invention relates to short interfering nucleic acids (siNA) which to modulate expression of EGFR genes in tissue explants

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RESULT 29
ADM36242/c
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Best Local S
Matches 14
This invention relates to novel membrane proteins that are associated with differentiation and/or metabolic function of adipocytes, in particular of mouse origin. Specifically, it refers to the isolated nucleic acid molecules encoding all or part of these proteins, appropriate antibodies and screening assays useful for the development or drug compositions derived thereof. The present invention describes these compositions as useful for the treatment of diseases associated with abnormalities of adipocyte function, such that they can be used to prevent, treat or diagnose obesity, hypertension, hyperlipaemia, diabeter and arteriosclerotic, antilipaemic and hypotensive activities. This oligonucleotide is a 5' RACE gene specific PCR primer used to amplify murine adipocyte-derived cDNA of the invention.
                                                                                                                                                                                                                                                                                                                                             Example 3; SEQ ID NO 37; 195pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                    polynucleotides encoding them for screening
                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse membrane and secretory proteins of adipocyte origin and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matsuzawa Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-JUL-2002; 2002JP-00201856
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-JUL-2003; 2003WO-JP008690
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     treating a wide range of cancers such as breast and ovarian cancer. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents a chemically modified siRNA targeted to the human HERI (EGFR) mRNA transcript.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO2004007711-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     obesity; hypertension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5' RACE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (TAKE ) TAKEDA CHEM IND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   anorectic; antidiabetic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCR; ss; mouse; murine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  03-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              for the treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      244 TCAGACATCCTGAA 257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                              diabetes, arteriosclerosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TCAGACATCCTGAA 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               primer used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Funahashi T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     γģ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; 2 C; 8 G; 2 T; 6 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         by ex vivo gene therapy), or in grafts and transplants of a variety of conditions. They may be used for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hyperlipaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         differentiation; metabolic function; adipocyte;
hyperlipaemia; diabetes; arteriosclerosis;
; antiarteriosclerotic; antilipaemic; hypotensiv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21
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Pred. No.
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              cc library by providing several mRNA, synthesising first and second cDNA cc strands from mRNAs, ligating a first adaptor to the 5' end and a second cc adaptor to the 3' end of the double stranded cDNA to produce an adaptor-cc modified cDNA, cleaving the adaptor-modified cDNA with restriction ce endonuclease Sfil to produce first a nucleic acid and cloning the first nucleic acid into an expression vector. Also described are: a cellular cc library (I) comprising the retrovital vector library produced by (M1), where the first nucleic acid is fused with 5' end of the second nucleic acid; and screening (M2) for a bioactive agent capable of altering a cellular phenotype involving introducing a retrovital expression vector library, produced by (M1) into several cells, screening several cells for a cell with an altered phenotype. (I) is useful for creating novel cell lines from cancer patients, for identifying modulators of immune response, in neurobiology for screening anti-spoptotics for preservation of finance and processing anti-spoptotics for preservation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Producing expression vector library by synthesizing cDNA fr
ligating with adaptor, cleaving adaptor modified cDNA with
to provide first nucleic acid and cloning the nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (SHEN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     retroviral vector library; cellular phenotype; cancer; immune i neurobiology; anti-apoptotic; neuronal function; neuronal death exocytosis; intracellular calcium cycling; sarcolemmal calcium cardiomyocyte; arrhythmia; bone morphogenic protein; hormone mi cardiomyocyte; arrhythmia; bone morphogenic protein;
                                                                                                                                                                                                                                                                                                                                                                                                                      The invention describes a nethod of producing an expression vector
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 6; Fig 1; 41pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Shen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08-MAY-2002; 2002US-00142662.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-MAY-2002; 2002US-00142662
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-NOV-2003
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YU S.
WU X.
PAYAN D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TGCACCTGCTGCTT 41
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20. .22
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/note= "This sequence overhangs the complementary stand
forming a sticky end"
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and prevention of neuronal
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      with vector.
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RESULT 31
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type B
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                                                                                                                                                          A CDNA fragment (see AAT10327) coding for the lambda light chain variable region of a human antibody which specifically recognises a lipopolysaccharide (LPS) antigen from Potential actual actual was cloned by PCR. The coding sequence can be used for preparing cell lines which produce monoclonal antibodies against the LPS. The present sequence is that of a 5'-primer used in the PCR amplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-JUN-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; immunoglobulin; IgM; lambda light chain; variable region; lipopolysaccharide; LPS; antigen; Pseudomonas aeruginosa; type monoclonal antibody; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-JUL-1996
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                                                                                                                                                                                                                                                                                                                      Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Anti-P.aeruginosa type B antibody VL region N-terminal 5'-primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (MITK ) MITSUI TOATSU CHEM INC
                            Local Similarity es 14; Conserv
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                                                                                                                                                                                                                                                                                                                                                                encoding human antibody variable regions against P.aeruginosa useful for prodn. of monoclonal antibody-producing cells.
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                                                                                                                                                                                                                                                                                                                    2; Page 11; 12pp; Japanese
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llarity 100.0%;
Conservative (
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                                                 2.8%;
                                                                                                                    9 C; 5 G; 4 T; 0 U; 0 Other;
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                                                   Score 14;
Pred. No.
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                                                                        DB 2;
                                                   1.3e+04;
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                                                                        Length 23;
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Query Match Best Local Similarity

2.8**%**; 78.6**%**;

Score 14; Pred. No.

DB 10; Length 1.3e+04;

23;

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RESULT 32
ADG29798
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11-MAR-2002; 2002US-0365782P.
06-JUN-2002; 2002US-0366782P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
15-JAN-2003; 2003US-0440129P.
                                        nootropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary disease such as restenosis, atherosclerosis, Alzheimer's disease or parkinson's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Furthermore, the siNA may be utilised for gene therapy applications. The current sequence is that of the siNA DNA-
                                                                                                                                                                                                                                                                                                                                                                                                                                       Mcswiggen
Jamison S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              double-stranded short interfering nucleic acid; siNA; antiarteriosclerotic; neuroprotective; nootropic; antiparkinsonia antiarteriosclerotic; neuroprotective; notropic; antiparkinsonia anticonvulsant; pulmonary disease; restenosis; atherosclerosis; Alzheimer's; Parkinson's; epilepsy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; ss; DNA-RNA hybrid;
                                                                                                                                                                                        target gene comprising one or more chemical modifications and eac of the double-stranded siNA comprises about 21 nucleotides. The sthe invention demonstrates antiarteriosclerotic, neuroprotective,
                            Parkinson's disease, epilepsy, amyotrophic lateral sclerosis. amyotrophic lateral sclerosis. The gene therapy applications. The RNA hybrid of the invention.
                                                                                                                                                                                                                          The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand
                                                                                                                                                                                                                                                                                                                                      New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene of for treating diseases that respond to modulation of gene expression or
                                                                                                                                                                                                                                                                                          Example 24;
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BP;
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Other;
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Matches
 11-MAR-2002;
11-MAR-2002;
29-MAY-2002;
06-JUN-2002;
06-JUN-2002;
03-JUL-2002;
03-JUL-2002;
29-AUG-2002;
09-SEP-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cancer; cytostatic; human; oncogene; epidermal growth factor receptor; EGFR; HER1; c-erb-B-1; DNA-RNA hybrid; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification;
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                                               2002US-0358580P.
2002US-0363124P.
2002WG-US016840.
2002US-00163552.
2002US-0386782P.
2002US-0393924P.
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   2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
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/mod_base= OTHER
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note= "2'-deoxy-2'-fluoro bases"
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e= "2'-deoxy-2'-fluoro
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 RESULT 34
ADO25209
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             interfering RNA (sirNA), double-stranded RNA, micro-RNA (mirNA) and short chairpin RNA (sirNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of EGFR genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating a wide range of cancers such as breast and ovarian cancer. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single mucleotide polymorphisms). The present sequence represents a chemically modified siRNA targeted to the human HERI (EGFR) mRNA transcript.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local
19-NOV-2002; 2002FI-00002064
                              19-NOV-2003; 2003WO-FI000888
                                                                                                                                                                              Bacterial species identification; bacterial infection; hyper-variable region; conserved region; topoisomerase
                                                                                                                                                                                                                            N. gonorrhoeae gyrß gene, oligonucleotide probe
                                                                                                                              Neisseria gonorrhoeae
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                                                                                                                                                                                                                                                                                              AD025209;
                                                                                                                                                                                                                                                                                                                             ADO25209 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        be double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to short interfering nucleic acids (siNA) which
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-697612/66.
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21-OCT-2002;
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                                                                                                                                                             Infection-causing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       interference.
                                                                                                                                                                                                                                                                                                                                                                                                                                          244 ТСАСАСАТССТСАА 257
                                                                                                                                                                                                                                                                                                                                                                                                            5 UCAGACAUCCUGAA 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ate expression of one or more human epidermal growth factor (EGFR) genes (including HER1, HER2 HER3 and HER4) by RNA nce. The siNNa may or may not comprise ribonucleotides and may or single stranded. They further comprise sense and antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        <u>.</u>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 6 A; 8 C; 2 G;
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2002US-00277494
2003US-0440129P
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                                                                                                                                                              jion; conserved region; topoisomerase; gyrB; parE;
bacteria; respiratory tract infection; probe; ss.
                                                                                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                              entry)
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .g. for treatment and
n of the epidermal growth
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Jamison
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RESULT 35
ADO25212
       The present invention relates to a diagnostic method for detecting and clidentifying bacterial species that cause infections. The method comprises contacting amplified DNA from a clinical sample with a desired combination of oligonuclectide probe sequences that hybridise under coronal hybridisation conditions with hyper-variable regions situated near the conserved regions of genes encoding topolsomerases, especially cyrlyparE, of bacterial species causing the infections. The invention causing bacteria, and a confidence of infection causing bacteria, and a convention. The combination of oligonuclectide probes and primers of the cinvention. The combination of oligonuclectide probes and primers of the species, such as those causing respiratory tract infections. The primers comprised are useful for amplifying topolsomerase genes especially genes encoding cyrls and parE proteins. The present sequence represents an analysis of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Diagnostic method for detecting and identifying bacterial species causing infections from clinical sample by contacting amplified DNA with a desired combination of oligonucleotide probe sequences.
    Diagnostic method for detecting infections from clinical sample
                                                                                                                                                         19-NOV-2003;
                                                                                                                                                                                                                                          Neisseria gonorrhoeae.
                                                                                                                                                                                                                                                                                  Bacterial species identification; bacterial infection; hyper-variable region; conserved region; topoisomerase
                                                                                                                                                                                                                                                                                                                         N. gonorrhoeae gyrB gene, oligonucleotide probe
                                                                                                                                                                                                                                                                                                                                                     26-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                 ADO25212;
                                                                                                                                                                                                                                                                                                                                                                                                            ADO25212 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   oligonucleotide probe of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim
                                             WPI; 2004-420639/39.
                                                                                                                              19-NOV-2002; 2002FI-00002064.
                                                                                                                                                                                                              WO2004046379-A1
                                                                                                                                                                                                                                                                   infection-causing
                                                                                                    (MOBI-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14;
                                                                                                    MOBIDIAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
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                                                                      Jalava J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 30; 77pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                         2003WO-FI000888
                                                                                                                                                                                                                                                                                                                                                     (first
                                                                                                                                                                                                                                                                                region; conserved
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11
                                                                                                                                                                                                                                                                   gion; conserved region; topoisomerase;
bacteria; respiratory tract infection;
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                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
                                                                         Nikkari
                                                                                                                                                                                                                                                                                                                                                     entry)
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and identifying bacterial species causing by contacting amplified DNA with a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.3e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 23;
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                                                                                                                                                                                                                                                                   parE;
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RESULT 36
AAQ78839/c
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the conserved regions of genes encoding topoisomerases, especially gyrB/parB, of bacterial species causing the infections. The invention also discloses broad-range primers originating from the conserved regions of the topoisomerase genes of infection-causing bacteria, and a diagnostic kit containing the oligonucleotide probes and primers of the invention. The combination of oligonucleotide probes is useful for detecting, identifying or classifying infection-causing bacterial species, such as those causing respiratory tract infections. The primers are useful for amplifying topoisomerase genes especially genes encoding gyrB and parE proteins. The present sequence represents an oligonucleotide probe of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to a diagnostic method for detecting an identifying bacterial species that cause infections. The method compx contacting amplified DNA from a clinical sample with a desired combination of oligonucleotide probe sequences that hybridise under normal hybridisation conditions with hyper-variable regions situated normal hyper-variable regions normal hyper-variable regions normal hyper-variable regions norma
                                                                                                                                                                                                                  Human monoclonal antibodies nucleic acids and cell lines
                                                                                                                                                                                                                                                                                        WPI; 1994-358194/44
                                                                                                                                                                                                                                                                                                                                        Burton
                                                                                                                                                                                                                                                                                                                                                                                                                                    30-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-NOV-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9425490-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             immunoassay;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human cytomegalovirus; antibody lambda light chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HCMV Ab
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19-JUL-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              desired combination of oligonucleotide probe sequences
                                                                                                                                                                                                                                                                                                                                                                                       (SCRI)
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                                                                                                                                                                                                                                                                                                                                        DR,
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                                                                                                                                                                                                                                                                                                                                     Barbas C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 11 A; 7 C; 2 G; 3 T; 0
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      chain variable domain PCR primer V(L7).
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                                                                                                                                                                                                               (MAbs) against producing the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 14;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                        Williamson
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hes 0;
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                                                                                                                                                                                                             human
MAbs,
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                                                                                                                                                                                                               cytomegalovirus useful in diagno
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AAQ78833-Q78839 are 5' primers which separately pair with the 3' prime AAQ78840, for the PCR amplification of the human cycomegalovirus (HMCY antibody lambda 11ght chain variable regions. The complete antibodies could be used to detect HCMV, and anti-human HCMV Abs in human patient via a competitive immunoassay. The antibodies may also be useful in

patients

(HMCV)

Example

2b1; Page 84; 171pp;

English

acids and cell

lines

diagnosis

also is and

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                                                                                                Query Match
Best Local S
Matches 14
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                       This invention relates to novel oligonucleotide PCR primers which may be used to identify markers associated with genetic diseases and traits in dogs, in particular to diagnose genetic diseases that are not phenotypically visible and to identify carriers of recessive diseases. A specific application is diagnosis of copper toxicosis (CT). The invention can also be used to create a genetic map of the canine genome; to generate breed-specific profiles; to establish patternity and to identify dogs from DNA fingerprints. The method provides rapid analysis of the target sequences from only a small sample of DNA. Diagnosis can be done at any time in the dog's life. The present sequence is that of a PCR
                                                                                                                                                                                                            target sequences from on
at any time in the dog's
primer of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New oligonucleotide primers for diagnosis of genetic diseases and traits in dogs - amplify specific regions of the genome containing microsatellite repeats, especially for diagnosing copper toxicosis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         genetic disease; genetic trait; dog; carrier of recessive disease; copper toxicosis; CT; canine genome map; breed-specific profile; DNA fingerprint; dog identification; PCR; primex; ss.
                                                                                                                                                                      Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-FEB-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Canine disease marker-related PCR primer 470
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                                                          GAAGAATTACCTGG 268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        standard;
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                         GAAGAATTACCTGG
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                                                                                                Conservative
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                                                                                                                                                                      A; 4 C; 4
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100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                          English
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                                                                                                               Score 14; DB 2; L
Pred. No. 1.3e+04;
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Pred. No. 1.3e+04;
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                                                                                                                                                                      T; 0 U; 0
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                                                                                                                                 Length 24;
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 RESULT 39
AAZ60776/c
ID AAZ607
XX
AAC AAZ607
XX
DT 16-MAY
DX 5' PCR
XX
CONfOR
XW Variab
KW Variab
KW Liver
KW PCR pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 38
AAV68558/c
ID AAV685
XX AAV685
XX Lambda
XX IAMBDIII
XX ARM AUTO-II
XX II
X
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Best Local S
Matches 14
               Variable heavy chain; human anti-HCV antibody; chronic Hepatitis C; conformation-dependent epitope; HCV glycoprotein E2; HCV infection; liver transplantation; Tupaia-hepatocyte; HCV-infectious human sera
conformation-dependent liver transplantation; PCR primer; ss.
                                                                                                                                                                                                            AAZ60776 standard;
                                                                                                                                   16-MAY-2000
                                                                                                                                                                      AAZ60776;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             targeting
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9846645-A2.
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(RAUM/) RAUM T.
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                                                                                            primer
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Sequence 24 BP; 5 A; 10 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This is the nucleotide sequence of a lambda chain fragment used for amplification in the method of the invention, for providing receptors that can be used for targeting antigens in humans without being immunogenic themselves. Such receptors can be used for treating diseases such as tumours or auto-immune diseases, graft rejection after transplantation, infectious diseases by targeting cellular receptors as well as allergic, inflammatory, endocrine and degenerative diseases by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Production of anti-human antigen receptors - by selecting a combination of functionally rearranged VH and VL immunoglobulin chains expressed from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Page 28; 84pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  a recombinant vector.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1998-594564/50.
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                                                                                                                                                                                                                                           GTGAGCTCAACCTG
                                                                                                                                                                                                                                                                                 GTGAGCTCAACCTG 198
                                                                                                                                                                                                                                                                                                                                                                                                                                                         key molecules involved in
                                                                                                                                                                                                                                                                                                                              Conservative
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                                            (first entry)
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for light
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5' PCR
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  chain variable region
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Pred. No.
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RESULT 40
ABZ30677
ID ABZ30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PCR primers AAZ60770-78 were used to amplify DNA encoding the variable light chain of a human anti-Hepatitis C virus (HCV) antibody. The amplified sequence was used to construct antibodies of the invention.
These antibodies comprise at least one complementarity determining region (CDR) of the variable domain of a human antibody that specifically recognizes a conformation-dependent epitope of HCV glycoprotein E2 and is capable of precipitating covalently or non-covalently associated E2/E1 complexes. The antibodies are useful for preventing (re) infection of HCV and are useful for alleviating chronic Hepatitis C in a human or an animal. The antibodies are useful for diagnosing chronic Hepatitis C for the presence of neutralization of binding of HCV glycoprotein E2 on to tablet cells. The antibodies are useful for the treatment or prevention of HCV infection or reoccurrence of HCV infection, in which they are administered prior, during or after liver transplantation. The antibody is also useful in liver transplantation. The antibody is also useful for the prevention of infection of Tupaia-hepatocyte with HCV-infectious human sera
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                       Fungus; yeast; tetracyclin; promoter; GRACE strain; biosynthesis; signal transduction; DNA replication; cell division; growth; proliferation; Candida albicans; fungicide; antifungal; PCR; primer;
     11-JUL-2002
                                                                                                                                                                                                                                                                       Candida albicans
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Virus, comprising at least one compl variable domain of a human antibody.
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                                                                                                             albicans
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                                                                                                                                                                                                                                                                       GRACE strain
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at least c
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100.0%; Pred. No. 1.3e+04;
Live 0; Mismatches 0;
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                                                                                                                                                                                                                                                                       PCR
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Mouse;

11-JUL-2002 AAL42621;

(first entry

tumour

gastrointestinal motility; post-operative ileus; central nervous system disease; migraine; CNS tr sleep disorders; acute pain; musculoskeletal pai

CNS trauma; schizophrenia;

septic shock;

post

pain;

peroxisome proliferator activated receptor-delta activator; PPAR-delta activator; inflammatory condition; shock state; haemorrhagic shock; traumatic shock; immune disorder;

Mouse inhibitor of nitric oxide synthase (INOS) antisense PCR primer.

PCR; primer; ss; inhibitor of nitric oxide synthase; necrosis factor TNF;

RESULT 41

AAL42621 standard; DNA; 25

밁 S

11

TTTCTGCTGTCTGA

24

334 TTTCTGCTGTCTGA 347

Query Match Best Local S Matches 14

Similarity

100.0%; 2,

Pred.

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1.3e+04; thes 0;

Indels

0;

0

Mismatches

Score 14; DB 6; Length 24;

Conservative

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Cells in which both alleles of a gene are modified, comprising modifying CC one allele by insertion or replacement by a cassette having an CC expressible selectable marker and modifying other allele by recombination, of a promoter replacement fragment with a heterologous CC promoter, so that expression of the second allele is regulated by the CC promoter. (M1) is useful for constructing a strain of diploid fungal CC cells in which both alleles of a gene are modified. The diploid fungal CC cells in which both alleles modified are useful for identifying a gene that CC contributes to the survival or growth of a fungus, a gene that CC contributes to the resistance of a diploid fungus to an antifungal agent that inhibits the growth of a fungus to an antifungal CC agent, an antifungal agent that inhibits the growth of a diploid fungus CC and for identifying a therapeutic agent for treatment of a mammalian CC disease. (M1) is useful for identifying a compound which modulates the CC compound catabolism, biosynthetic, transporter, transcriptional, CC translational, signal transduction, DNA replication and cell division CC ability to inhibit growth or proliferation of C. abicans cells and for treating infection by C. albicans. The present sequence is that of a PCR CC primer used in the method of the invention. Note: The sequence data for CC this patent is not represented in the printed specification but is based CC on sequence information supplied to Derwent by the European Patent Office
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Constructing strains for identifying gene products as effective targets for therapeutic intervention, by inactivating in the strain one allele of a gene and placing other allele of the gene under conditional expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-DEC-2000; 2000US-0259128P
20-FEB-2001; 2001US-00792024
22-AUG-2001; 2001US-0314050P
    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 36; SEQ ID NO 4828; 167pp + Sequence Listing; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-566694/60.
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    BP;
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modifying
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PN WO2
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                        inflammatory bowel disease; adult respiratory distress syndrome; eczema; cystic fibrosis; chronic obstructive pulmonary disease; dermatitis; multiple solerosis; glomerulonephritis; psoriasis; urticaria; asthma; glaucoma; systemic lupus erythematosis; sepsis; atheroselerosis; hypoxic; ischaemic heart disease; nitric oxide synthase inhibitor; iNOS; ss.
                                                    Mus sp
                                                                                                                                                             Mouse; PCR; primer; murine; antiinflammatory; inflammatory disease; peroxisome proliferator activated receptor-delta; pphk delta activator; joint; rheumatoid arthritis; osteoarthritis; prosthetic joint failure; gastrointestinal tract; ulcerative colitis; Crohn's disease; burn injur;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention comprises a method for treating diseases for which an inhibitor of nitric oxide synthase (INOS) and/or tumour necrosis factor (TNP) is indicated. The method of the invention involves administering a peroxisome proliferator activated receptor-delta (PPAR-delta) activator. The method of the invention is useful in the treatment of: inflammatory conditions; shock states (e.g. septic shock, haemorrhagic shock or traumatic shock); immune disorders; disorders of gastrointestinal motility (e.g. post-operative ileus); diseases of the central nervous system (e.g. migraine, CNS trauma, schizophrenia or sleep disorders); acute pain (e.g. musculoskeletal, post operative, or cancer); and HIV infection. The present DNA sequence represents a mouse inhibitor of
                                                                                                                                                                                                                                                       Mouse INOS anti-sense PCR primer.
                                                                                                                                                                                                                                                                                             11-JUL-2002
                                                                                                                                                                                                                                                                                                                             AAK99746;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 25 BP; 2 A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 22; 30pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       synthase (INOS)
                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6 C; 10 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 14; DB; Pred. No. 1.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 6; LC
0, 1.3e+04;
0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ٥,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                injury;
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WO200228434-A2

16-MAR-2001; 2001US-0276759P

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RESULT 43
ACK25054/c
ID ACK250
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        comprises the administration of a peroxisome proliferator activated creceptor-delta (PPAR delta) activator. The peroxisome proliferator can be used in the treatment of inflammatory diseases or conditions, e.g. of joints creatment of inflammatory diseases or conditions, e.g. of joints consistent of inflammatory diseases or conditions, c.g. of joints consistent of inflammatory diseases or conditions, consistent of inflammatory collits, Crohn's disease, inflammatory collits, Crohn's disease, athma, c.g. of joints consistent of inflammatory disease, inflammatory collits, crohn's disease, asthma, c.g. (multiple sclerosis), pancreas, kidneys (glomerulonephritis), skin c.g. (dermatitis, psoriasis, eczema, urticaria or burn injury), eyes c.g. (dermatitis, psoriasis, eczema, urticaria or burn injury), eyes c.g. (glaucoma), multi-organ diseases (systemic lupus erythematosis or c.g. (segsis), inflammation of viral or bacterial infections atherosclerosis or hypoxic or ischaemic heart disease. This polynucleotide sequence crepresents a PCR primer used in reverse transcriptase amplification of investigating the inhibition of iNOS activity by PPAR delta activators of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                          15-MAR-2002; 2002US-00098263
                                                                                                                                                                                              EST; ss; probe; expressed sec
genetic variation; biallelic
                                                                                                               US2003104410-A1
                                                                                                                                                                              cross-species comparison.
                                                                                                                                                                                                                                              Human microarray DNA oligonucleotide SEQ ID
                                                                                                                                                                                                                                                                                                                                                  ACK25054 standard; DNA; 25
                                                                                                                                                                                                                                                                               14-OCT-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 25 BP; 2 A; 6 C; 10 G; 7 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Method, useful in the treatment of inflammatory diseases or conditions, e.g. arthritis, asthma, multiple sclerosis or dermatitis, comprises administration of a peroxisome proliferator activated receptor-delta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a method for treating inflammatory diseases, comprises the administration of a peroxisome proliferator activated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 2; Page 21; 29pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Buchan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-394318/42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-OCT-2000; 2000GB-00024361
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-OCT-2001; 2001WO-GB004373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GLAX ) GLAXO GROUP LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity nes 14; Conserv
                                                                                                                                              sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   305
                                                                                                                                                                                                                                                                                                                                                                                                                                     æ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AGAGCCTCGTGGCT 318
                                                                                                                                                                                                                                                                                                                                                                                                                                  AGAGCCTCGTGGCT 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.8%; Scilarity 100.0%; F
                                                                                                                                                                                                            sequence tag; microarray;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 14; DB; Pred. No. 1.3
                                                                                                                                                                                           marker; polymorphism; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.3e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6
                                                                                                                                                                                                                                            NO 125035
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                           gene expression;
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RESULT 44
ACH63234
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              of at least one target sequence. The method of analysis comprises comprises to hybridising at least one or more nucleic acids to at least two or more concluded and probes and detecting the hybridisation. The nucleic acid probes are attached to a solid support. The analysis comprises monitoring concludes are attached to a solid support. The analysis comprises monitoring concludes are attached to a solid support. The analysis comprises monitoring concludes acid nevels, identifying biallelic markers or polymorphisms, concluded acids further comprises a tag sequence. The array of nucleic acid probes is useful in in situ hybridisation, in Southern, Northern or dotton to identify or detect the sequence or specific mutations of any gene, in mapping the 5' termini of mRNA molecules by primer extensions or in screening cDNA or genomic libraries or subclones contained and previously sequenced. The sequence presented is one of the nucleic acid probes incorporated in the microarray. Note: The sequence concluded and previously sequenced in the microarray. Note: The sequence concluded and previously sequenced in the microarray. Note: The sequence concluded and previously sequenced in the microarray. Note: The sequence concludes and so be obtained in electronic format directly concluded and sequence and so be obtained in electronic format directly concluded and sequence and so be obtained in electronic format directly concluded and sequence and sequence and sequence are sequenced.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention discloses a microarray comprising a plurality of nucleic acid probes including one of 2,018,500 fully defined sequences, or its perfect match, perfect mismatch, antisense match or antisense mismatch. Also disclosed is a method of gene expression analysis. The array is used in monitoring gene expression levels by hybridisation to a DNA library, in analysis of genetic variation or in hybridisation of tag-labelled compounds. The nucleic acid probes are specifically designed for analysis of at least one target sequence. The method of analysis comprises
                                                                                                                                                                                                                             Gene expression analysis; array; hybridisation; genetic variation; tag-labelled compound; gene family; in situ hybridisation; library screening; Southern hybridisation; northern hybridisation; dot-blot hybridisation; gene sequence; mutation detection; target sequence; probe; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New array of nucleic acid probes, useful for Southern, Northern or dot-blot hybridization
                 08-AUG-2001; 2001US-0311040P
                                                            08-AUG-2002; 2002US-00215112
                                                                                                                                                                                                                                                                                                                                                                 DNA target
                                                                                                                                                                                                                                                                                                                                                                                                            17-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ACH63234 standard; DNA; 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; SEQ ID NO 125035; 9pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-567953/53
                                                                                                       01-MAY-2003
                                                                                                                                                 US2003082596-A1
                                                                                                                                                                                            Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (AFFY-) AFFYMETRIX INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  for this patent can also be obtained in USPTO at segdata.uspto.goc/sequence.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CCGACCGGAGAGCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CCGACCGGAGAGCG 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    or specific mutations of any gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A.
                                                                                                                                                                                                                                                                                                                                                             #12370 useful in array
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9 C; 8 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        w
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               股
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 14; DB 9; L
Pred. No. 1.3e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                             for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           t i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         <u>..</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 25;
                                                                                                                                                                                                                                                                                                                                                             genetic analyses.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         situ hybridization, identify or detect (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           , in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0,
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The present invention relates to nucleic acid sequences that are CC complementary to particular genes, and can be used as probes for a CC variety of analyses such as gene expression analysis. Each probe CC comprises 9 or more consecutive nucleotides from at least one of 14336 comprises 9 or more consecutive nucleotides from at least one of 14336 concleotide sequences defined in the patent, or their perfect sense match, CC sense mismatch, antisense match or antisense mismatch oligonucleotides. The probes may be used in an array comprising at least 10 distinct concleic acid probes. The array is useful in monitoring gene expression CC levels by hybridisation to a DNA library, in analysing genetic consecutions, and in hybridising tag-labelled compounds. The probes are useful for identifying family members of a gene. The probes are also useful in in situ hybridisations, in screening cDNA or genomic libraries (or derived subclones) for additional clones containing segments of DNA CC that have been previously isolated and sequenced, in Southern, northern, cc or dot-blot hybridisation of genomic DNA to identify or detect the CC sequence of any gene or detect specific mutations in any gene, and in CC sequence of any gene or detect specific mutations in any gene, and in CC sequence of the invention are also useful as PCR primers. CC complementary to particular genes with a wide range of analytical uses. CC The invention provides a large collection of nucleic acid sequences cC complementary to particular genes with a wide range of analytical uses. CC The sequence data for this patent was obtained in electronic format the target sequences of the invention. Note: The sequence data for this patent was obtained in electronic format complements.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New probe array useful e.g. for monitoring gene expression levels, for analyzing genetic variations, or for hybridizing tag-labeled compounds, comprises multiple nucleic acid probes.
   Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-576608/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; SEQ ID NO 12370; 9pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (MITT/) MITTMANN
BP; 4 A; 9 C; 7 G; 5 T; 0 U; 0 Other;
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밁
                                Matches
                                        Query Match
Best Local
                126 CTGCCACCTGTACT 139
                                 14;
s
                                         Similarity
CTGCCACCTGTACT 18
                                 Conservative
                                        100.0%;
                                 0
                                         Score 14;
Pred. No.
                                 Mismatches
                                         1.3e+04
                                                DB 9;
                                 ٥.
                                                Length 25
                                 Indels
                                 0
                                 Gaps
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RESULT 45
ABV74186
  Murine
                                                                                ABV74186 standard; DNA; 26
                            27-JAN-2003
                                                      ABV74186;
FIZZ1 probe mFIZZ1.probe1.
                          (first entry
                                                                                ВP
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obesity; anorectic; gene therapy; probe; ss. FIZZ1; mFIZZ1; mouse; found in inflammatory zone-1; metabolic disorder;

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(GETH ) GENENTECH INC
            30-MAR-2001; 2001US-0280571P
29-MAR-2002; 2002US-00112917
                          29-MAR-2002; 2002WO-US010136
                                   10-OCT-2002
                                                     Mus musculus
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RESULT 46
AAC63911/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local
                                                                                     Resolving 2-oxobicyclo(3.1.0)hexane-6-carboxylates into enantiomer esters and enantiomer acids useful as pharmaceutical intermediates by contacting enzyme with 2-oxobicyclo(3.1.0)hexane-6-carboxylate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Esterase; enantiomer resolution; racemic mixture; 2-oxobicyclo(3.1.0)hexane-6-carboxylate ester; pharmaceutical synthetic intermediate; plasmid construction; mutagenic oligonucleotide; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-AUG-2003
09-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antagonist activity of a compound, to screen a subject for a FIZZI related metabolic disorder, to measure the obesity-reducing activity of a modality, to reduce the metabolic activity of a subject, and to alter expression of FIZZI in the WAT of a subject. The FIZZI nucleic acids and proteins are useful in the treatment of metabolic disorders such as obesity, cachexia and increased metabolic rate caused by severe burns
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence is that of a probe, designated mFIZZ1.probel, which was used in an example of the invention for quantitation of FIZZ1 (found in inflammatory zone-1) mRNA in mice. Results showed that FIZZ1 expression is markedly depressed in the white adipose tissue of obese, leptin-deficient ob/ob mice. The invention provides methods of using human and murine FIZZ1 polynucleotides, polypeptides and antibodies to increase metabolic activity in a subject, to measure the FIZZ1 agonist of the contract of
                                                                                                                                                                                                                                                                                                                                                          16-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                          17-APR-2000; 2000EP-00108383.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             25-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chromobacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chromobacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAC63911 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Increasing metabolic activity in a e.g., obesity comprises increasing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-040672/03
                                                                                                                                                                                                                                                                                                       (SUMO)
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                                                                                                                                                                                                     2001-000879/01.
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                                                                                                                                                                                                                                                                                                       SUMITOMO CHEM CO LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                     Takashima Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TGACTGCTACTGGG 487
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TGACTGCTACTGGG 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 3 A; 4 C; 11 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                       99JP-00109645
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SC-YM-1 esterase mutation/PCR primer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.8%;
                                                                                                                                                                                                                                                     Mine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               subject for treating or activity of FIZZ1 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.3e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ID NO:10
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Example; Page

invention

relates 29; 35pp;

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resolving enantiomers

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Sequence

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0 Other;

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English. methods of

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RESULT 47
AAT34531
ID AAT34
XX AAT34
XX AAT34
XX IOO
DE Human
XX Fas a
KW Fas a
KW Pas a
KW Pas a
KW O962
XX O04-JU

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                                                 A PCR primer (AAT34531) is based on nucleotides 1-22 of human Fas ant CDNA (see also AAT34536). It was used with a primer (AAT34536) complementary to nucleotides 1316-1336 of the cDNA to amplify human f mRNA from nucleotides 170-1336. The template mRNA was obtained from typeripheral blood mononuclear cells of healthy subjects and from systelupus erythematosus (SLE) and angioimmunoblastic lymphadenopathy patients. PCR products were cloned into a pCR vector, expressed in E. coli, and sequenced (see also AAT34533-34). 4 Distinct mRNA variants also AAT34527-30) were identified
                                                                                                                                                                                                                                                                                                                                                                                       Natural, soluble form of Fas antigen secreted by human cells : alternative mRNA processing - used to diagnose Fas-associated e.g. systemic lupus erythematosus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  oxobicyclo(3.1.0) hexane-6-carboxylate esters such that one enantiomer is in an ester form and the other enantiomer is in an acid form. The methods involve contacting 2-oxobicyclo(3.1.0) hexane-6-carboxylate ester with an enzyme which preferentially hydrolyses one enantiomer of the ester of 2-oxobicyclo(3.1.0) hexane-6-carboxylate to obtain one enantiomer as an acid and the other as an ester. The methods are used to resolve 2-oxobicyclo(3.1.0) hexane-6-carboxylates, which are useful as intermediates for pharmaceuticals. An example of an enzyme which may be used in the methods of the invention is an esterase from Chromobacterium SC-YM-1 (FERM BP-6703 transferred from FERM P-14009). The present sequence represents an oligonucleotide used to generate a mutant esterase and which was also used as a PCR primer in plasmid construction. (Updated on 06-AUG-2003 to correct OS field.)
                                                                                                                                                                                                                                                                                                                                        Example 1; Page 74; 152pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAT34531;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     reaction;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cheng
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                                                                                                                                                                                                                                                                                                                                                                                                                    is result of disease,
                                                                                                                                                                                                                                                                             Fas antigen
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                                                                                                                                                                 systemic
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RESULT 48
AAT34533
ID AAT34
XX Pas angle o
XX Pas angle o
XX Polym
XX WO962
XX WO962
XX WO962
XX YOUNE
XX WO962
XX WAT1,
XX WAT1,
XX MATU,
PT Alte;
PT e.g.
XX Natu,
PT Alte;
PT e.g.
XX CC A pr.
CC CDNA
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Best Local :
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                                                                                                                                             Matches
                                                                                                                                                                                                  A primer (AAT34533) is based on nucleotides 377-3912 of human Fas antiger cDNA (see also AAT34526). It was used with a primer (AAT34534) complementary to nucleotides 1059-1076 of the cDNA to sequence PCR products amplified (see also AAT34531-32) from fas mRNA derived from the peripheral blood mononuclear cells of healthy subjects and from systemic lupus erythematosus (SLE) and angioimmunoblastic lymphadenopathy patients. 4 Distinct mRNA variants (see also AAT34527-30) were identified
                                                                                                                                                                                                                                                                                                 Natural, soluble form of Fas antigen secreted by human cells is alternative mRNA processing - used to diagnose Fas-associated die.g. systemic lupus erythematosus.
                                                                                                                                                                                 Sequence 15 BP; 4 A; 7 C; 3 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                         WPI; 1996-321796/32.
                                                                                                                                                                                                                                                                                                                                                            Mountz
                                                                                                                                                                                                                                                                                                                                                                                                                     22-DEC-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                polymerase chain reaction; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fas antigen; autoimmune disease; systemic lupus erythematosus; SLE; angioimmunoblastic lymphadenopathy; AILD; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human Fas antigen primer.
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                                                                                                                                                                                                                                                                               Example 1; Page 75; 152pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                  23-DEC-1994;
         15-SEP-2003
29-AUG-2000
                                      AAA38405;
                                                        AAA38405
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                                                                                                                                            13; Conserv
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                                                        standard;
                                                                                                                                                                                                                                                                                                                                                           Liu C,
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                                                                                                                                            Conservative
         (revised)
(first entry)
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                                                                                                                                                     2.6%;
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                                                         DNA; 17
                                                                                                                                                                                                                                                                                                                                                            Zhou
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                                                                                                                                                              DB 2;
                                                                                                                                                      4.2e+04;
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AAS12968
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Best Local S
Matches 13
                                                                                                                                                                                                                                                                               Sequence 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-SEP-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pseudomonas sp;
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         20-SEP-2001.
                             WO200168917-A1
                                               Homo sapiens.
                                                                  genetic
                                                                          Human; familial
                                                                                              Familial
                                                                                                                18-DEC-2001
                                                                                                                                    AAS12968
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Benzothiophene oxidase; bto gene
                                        WF505 bto gene cluster primer,
    cluster; desulphurisation; crude oil;
                                              SEQ
                                              ID NO:28
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98JP-00272744.

New proteins useful for desulfurization of crude oil have ability to control expression of benzothiophene oxidase group.

Example 1; Page 8; 43pp; Japanese.

encompasses genetic constructs and vectors comprising the genes encoding these proteins, hosts containing such constructs and the recombinant production of the enzymes. Pseudomonas sp. WF505 benzothiophene oxidase, benzothiophene oxidase co-enzyme and bto gene cluster regulator are useful in the desulfurisation of crude oil and petroleum products, and for the clarification of soil and waste water contaminated by organic sulphur compounds. Sequences AAA38401-A38423 represent primers used in the exemplification of the invention in the isolation and characterisation of the genes of the Pseudomonas sp. WF505 benzothiophene oxidase gene cluster. (Updated on 15-SEP-2003 to standardise OS field) benzothiophene oxidase co-enzyme (AAY97824), which controls benzothiophene activity via redox regulation. These enzymes are respectively encoded by the bto gene cluster open reading frames ORF2-16 (AAA38398), ORF4-9 (AAA38396) and ORF6-7 (AAA38395). The invention also The invention relates to proteins encoded by the benzothiophene oxidase (bto) gene cluster from Pseudomonas sp. WF505, specifically the protein which controls expression of the gene cluster (AAY97827), the benzothiophene oxidase (bto) enzyme itself (AAY97825) and a WF505 benzothiophene ORF2-16

8 C; 3 G; 4 T; 0 U; 0 Other;

Conservative 2.6. 100.0%; FI Score 13; DB 3; Li Pred. No. 4.2e+04; Mismatches Length 17; Indels 0, Gaps 0;

AAS12968 standard; DNA; 17 BP.

(first entry)

dysautonomia disease gene marker 88B2GA PCR primer #1.

dysautonomia; chromosome 9q31-33; PCR primer; ss

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RESULT 51
ABN07848/c
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Best Local :
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21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
                                                                     30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a method for identification of an individual carrying a gene associated with familial dysautonomia. This involves detecting the presence of a polymorphism located between D9312 and D9310 inclusive on human chromosome 9, which is linked to the gene associated with familial dysautonomia. Hence the presence of the polymorphism is indicative of the presence of the disease-associated gene. A nucleic aci sequence comprising the familial dysautonomia gene, and PCR primers encoding sequences flanking markers of the gene are useful for carrying out genetic testing of the disorder. This sequence represents a PCR primer used to detect the 88B2GA marker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; hear muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Identifying an individual carrying a gene associated with familial dysautonomia, comprises detecting a polymorphism located between D9S12 dysautonomia, comprises detecting a polymorphism located between D9S12 dysautonomia, comprises detecting a polymorphism located between D9S12 distributions on human chromosome 9, linked to the gene associated distributions of the compression of the compre
                                                                                                                                                     30-JAN-2001;
                                                                                                                                                                                                                                                                                                          25-MAY-2001; 2001WO-US016981
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human GDMLP-1 17-mer scanning
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13; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CCTGGGTGACAAG 14
             ; 2000US-0207456P.
2000US-023458PP.
2000US-0236359P.
2000GB-00024263;
2001WO-US000661.
2001WO-US000663.
2001WO-US000663.
2001WO-US000665.
2001WO-US000665.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gusella JF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.6%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO:5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequence SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            hGDMLP-1; heart;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ID NO:7840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A nucleic acid
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RESULT 52
ABNO7850/c
ID ABNO78
XX ABNO78
AC ABNO78
XX 29-MAY
XX Unman;
XX Human;
XX Human;
XX Human;
XX Homo s
XX WO2001
XX WO2001
XX YX WO2001
XX YX YX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   문
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CC protein 1 (http://linkup.nl). The protein and polynucleotide sequences of hGDMLP-1 CC nucleic acids can be used as probes to detect, characterise and quantify CC hGDMLP-1 nucleic acids in samples, as amplification substrates, to CC provide initial substrates for the recombinant engineering of hGDMLP-1 CC protein variants having desired phenotypic improvements, and for CC expressing the proteins. The hGDMLP-1 proteins or polypeptides may be CC used as immunogens to raise antibodies that specifically recognise hGDMLP CC -1 proteins, as standards in assays used to determine the concentration CC and/or amount specifically of hGDMLP specific biomolecule CC capture probes for surface-enhanced laser desorption ionisation, as CC therapeutic supplement in patients having specific deficiency in hGDMLP concentration, and in vaccines or for replacement therapy. The CC polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a CC disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence in the expression of the present invention. N.B. CC may be used for the present of the present and the screening of the sequence data for this patent did not form part of the printed content of the 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local S
Matches 13
                                                                                                                                                                                                   muscle; m
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30-JAN-2001;
05-FEB-2001;
                                                                                                                                                                                                                       Human; genome-derived myosin-like protein 1; GDMLP-1; muscle; myosin; chromosome 22; gene therapy; vaccine;
                                                                                                                                                                                                                                                                                       Human GDMLP-1
                                                                                                                                                                                                                                                                                                                                  29-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                             ABN07850;
                                                                                                                                                                                                                                                                                                                                                                                                                        ABN07850 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 4 A; 3 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New polypeptide, for raising antibodies that recognize hgDMLP-1 protor as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           at ftp.wipo.int/pub/published_pct_sequence
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30-JAN-2001; 2001WO-US000670
30-JAN-2001; 2001WO-US000670
05-FEB-2001; 2001US-0266860P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 7840; 214pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-179446/23.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30 CACCIGCIGCITC 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13;
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                                                                                                                                                                                                     muscle
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                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                     17-mer scanning SEQ
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                                                                                                                                                                                                 chromosome
disorder; a
                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4
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                                                                                                                                                                                                 amplicon; screening; ss
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                                                                                                                                                                                                                                                                                  ID NO:5 sequence
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                                                                                                                                                                                                                                                                                     SEQ ID NO:7842
                                                                                                                                                                                                                       heart disease;
                                                                                                                                                                                                                                         hGDMLP-1; heart;
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25-MAY-2001; 2001WO-US016981

06-DEC-2001.

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RESULT 53
ABN07851/c
ID ABN078
XX
ABN078
XZ
AC ABN078
XZ
DT 29-MAY
XX
DT 29-MAY
XX
XX
Human
XX
KW Human;
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                                                                                                                                                                                                                                                                                                         The present invention describes a human genome-derived myosin-like CC protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 cc l can be used in gene therapy and vaccine production. The hGDMLP-1 cc mucleic acids can be used as probes to detect, characterise and quantify conclude initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for cexpressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP cused as immunogens to raise antibodies that specifically recognise hGDMLP concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser describing the concentration cand/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser describing in the capture of the present in patient in patients having specific deficiency in hGDMLP-1 concentration, as characteristic production, and in vaccines or for replacement therapy. The polymucleotide sequences encoding hGDMLP-1 may be used for diagnosing a concentration of horder associated with the expression of hGDMLP-1, in particular heart concentration of the present sequence and the expression of hGDMLP-1 is concentrated to chromosome 22.

The sequence data for this patent did not form part of the printed capture with published not sequence format directly from WIPO can the wine int/mulnitaged not sequence format directly from WIPO
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30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart;
                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gu Y,
                               Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:7843.
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                                                                                                         ABN07851 standard; DNA; 17
                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                   ftp.wipo.int/pub/published_pct_sequence
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                                                                                                                                                                                                                                                                            17
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                                                                                                                                                                         CACCTGCTGCTTC
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2001WO-US000669.
2001WO-US000670.
2001US-0266860P.
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2001WO-US000665.
2001WO-US000666.
2001WO-US000667.
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2000US-0234687P.
2000US-0236359P.
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2001WO-US000662.
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                                                        (first entry)
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                                                                                                                                                                                                                                                                           2 C; 8
                                                                                                                                                                                                                                     2.6%; Score 13; DB 6; Li
100.0%; Pred. No. 4.2e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              7842; 214pp; English
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                                                                                                           BP
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                                                                                                                                                                                                                                                                          G; 2 T; 0 U; 0 Other;
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Query Match Best Local Matches

Local

Similarity

100.0%;

Score 13; DB 6; Length 17; Pred. No. 4.2e+04;

4.2e+04; hes 0;

Indels

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Gaps

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Conservative

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13 30

CACCIGCIGCITC 1

CACCTGCTGCTTC 42

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CC protein 1 (hgpMLP-1). The protein and polyuncleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 conclete acids can be used as probes to detect, characterise and quantify composite the protein and substrates, to composite the protein substrates, to composite the protein substrates, to composite the protein variants having desired phenotypic improvements, and for composite the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The colymnicleotide sequences encoding hGDMLP-1 may be used for diagnosing a clistorider associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22.

The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present invention. N.B.

The sequence data for this patent did not form part of the printed captured to the printed control of the printed patent did not form part of the printed captured to the printed cap
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30-JAN-2001;
30-JAN-2001;
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21-SEP-2000;
27-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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   Sequence
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30-JAN-2001;
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2000US-0236359P.
2000GB-00024263.
2001WO-US000661.
2001WO-US000662.
2001WO-US000664.
2001WO-US000665.
2001WO-US000666.
2001WO-US000666.
2001WO-US000667.
2001WO-US000668.
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RESULT 54
ABNO7847/c
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XX DE Human;
XW Human;
XW Human;
XW muscle
XX W MO2001
XX Homo s
XX WO2001
XX O6-DEC
XX 26-MAY
PR 21-SEP
PR 04-OCT
PP 04-OCT
PR 30-JAN

                  CC The present invention describes a human genome-derived myosin-like CC protein 1 (hdDMLP-1). The protein and polynuclectide sequences of hdDMLP-1 can be used in gene therapy and vaccine production. The hdDMLP-1 calds can be used as probes to detect, characterise and quantify chorule initial substrates for the recombinant engineering of hdDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hdDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hdDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hdDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser description ionisation, as therapeutic supplement in patients having specific deficiency in hdDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hdDMLP-1 may be used for diagnosing a disorder associated with the expression of hdDMLP-1, in particular heart and skeletal muscle disorders. hdDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hdDMLP-1 sequence in the exemplification of the present invention. N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from MIPO at first part of the printed specification.
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21-SEP-2000;
27-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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2001WG-US000665.
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2001WG-US000666.
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2001WG-US000668.
2001WG-US000669.
2001WG-US000669.
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heart disease;
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RESULT 55
ABNO7849/c
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Best Local
The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as
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30-JAN-2001;
30-JAN-2001;
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04-OCT-2000;
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21-SEP-2000;
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RESULT 56
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06-JUN-2001;
10-SEP-2001;
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acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ5989 - ABZ62216, ABZ64544 - ABZ65531, ABZ66520 - ABZ66524, ABZ66530 - ABZ66595 represent substrate/target sequences for the human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                      Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
                                                                                                                                                                                                                                                                                                                                                                 Claim 58; Page 98; 185pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-140484/13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mcswiggen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABZ60565 standard; RNA;
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                                                                                                                                                                                                                                                                                                         The invention relates to a novel short interfering RNA (siRNA) nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-MAR-2003 (first entry
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNAzyme substrate
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2001US-0294140P.
2001US-0296249P.
2001US-0318471P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ω
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                                                                                                                                                                                                                                                                                                                                                                    English
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Pred. No. 4.2e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     #677
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Query Match Best Local

Similarity

2.6%;

Score 13; Pred. No.

DB 10;

Length 17;

Sequence

17

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Other;

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CC fragments of at least 15 consecutive nucleotides of these nucleotides, a CC sequence having at least 80% identity, after optimal alignment, with the nucleotides, a sequence that hybridizes under stringent conditions with the cc nucleotides. The nucleotides are used as probes or primers for detecting, cidentifying, quantifying and/or amplifying nucleic acids, as in vitro cc sense and antisense sequences, of nucleotides involved in tumour cs suppression or reversion, apoptosis and or viral resistance, to produce recombinant polypeptides, and to prepare transgenic animals, as cc experimental models. The nucleotides (also vectors containing them and cc ells containing the vectors), the encoded polypeptides and antibodies (Ab) against the polypeptide are useful for prevention and/or treatment co f viral infections or diseases characterized by development of tumours or cell degeneration (e.g. Alzheimer's disease or schizophrenia).

CC Analysis of the expression of the nucleotides can be used for diagnosis can and/or prognosis of these diseases. The nucleotides and polypeptides can algo be used to screen for their specific interactive molecules, corponentially useful for treating diseases associated with abnormal corponentially useful for treating diseases associated with abnormal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                В
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ADB45449/c
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Best Local S
Matches 13
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polypeptide
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                                                                                                                                                                                                                                                                                                                                                                                                                 New nucleic useful e.g.
                                                                                                                                                                                                                                                                                                                                                                 Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-441574/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Telerman A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-SEP-2001; 2001FR-00011981.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     virus resistance; transgenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tumour suppression/reversion associated nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADB45449;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADB45449 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (MOLE-) MOLECULAR ENGINES LAB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             195 CCTGGGTGACAAG 207
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    tic; antiviral; neuroprotective; nootropic;
probe; tumour suppression; tumour reversion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CCTGGGTGACAAG 3
                                                                                                                                                                                                                                                                                                                                                                                                               for treatment of tum
                                                                                                                                                                                                                                                                                                                                                               Page 706; 771pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                              and antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Amson R,
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                 tumors
                                                                                                                                                                                                                                                                                                                                                                                                                              prostate membrane-specific antigen,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                               and viral infection, also related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4.2e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      reversion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              <u>,</u>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   neuroleptic; ss;
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RESULT 59
AAV48442
ID AAV48
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AC AAV48
AC AAV48
CT 15-OC
XX
DE Trane
XX
KW Trane
KW modu:
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AAQ26547/c
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                                                                                                                                                                                                            Query Match
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 Transforming modulate; ger
                             Transforming growth factor beta-1 antisense oligonucleotide N30.
                                                                                                                                                                                                                                                  This probe was used together with AAQ26546-51 to analyse controls by oligonucleotide typing, whilst investigating polymorphisms - see AAQ26544,5
                                                    15-OCT-1998
                                                                                            AAV48442 standard; DNA; 18 BP
                                                                                                                                                                                                                                 Sequence 18
                                                                                                                                                                                                                                                                                              Example 2;
                                                                                                                                                                                                                                                                                                                       DNA encoding RING4, RING10, RING11 AND RING12 proteins - for treatment and diagnosis of immune disorders and screening of new immunosuppressants
                                                                                                                                                                                                                                                                                                                                                                                                                      19-DEC-1990;
16-SEP-1991;
                                                                                                                                                                                                                                                                                                                                                           WPI; 1992-250030/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-JUL-1992
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                                                                                                                                                                                                                                                                                                                                                                                Trowsdale
                                                                                                                                                                                                                                                                                                                                                                                                 (IMCR ) IMPERIAL CANCER RES TECHNOLOGY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             W09211289-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     polymerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MHC class I molecules;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Control probe #2 for caucosoid RING11 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        08-JAN-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ26547;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-DEC-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immunosuppressants; immunoenhancers;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .mmune disorders;
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                                                                                                                                                                                        1 Similarity
13; Conserv
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                                                                                                                                                                   CTGCCTGCAGCCT 123
                                                                                                                                                CTGCCTGCAGCCT 1
ing growth factor beta-1; TGF beta-1; antisense oligonucleotide;
gene expression; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                              'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     disease;
chain rea
                                                                                                                                                                                                                                                                                             Page 40;
                                                                                                                                                                                                                               BP; 4 A; 4 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                        Conservative
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                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                                              Kelly AP,
                                                                                                                                                                                                                                                                                                                                                                                                                      90GB-00027520.
91GB-00019711.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     91WO-GB002278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            transporter peptides; proteasome complex; ules; HLA; antigen processing; antigen presente; ankylosing spondylitus; prenatal diagnosis;
                                                                                                                                                                                                                                                                                          101pp;
                                                                                                                                                                                                2.6%;
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                                                                                                                                                                                                                                                                                            English.
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                                                                                                                                                                                                  Score 13;
Pred. No.
                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           treatment; diagnosis; screening;
                                                                                                                                                                                                           DB 2;
                                                                                                                                                                                                 4.2e+04;
                                                                                                                                                                                                          Length 18
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RESULT 60
AAZ77396/c
ID AAZ77
XX AAZ77
XX AAZ77
XX IO-SE
DT 10-SE
DX Human
XX Ampli

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Best Local S
Matches 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CC AAV48412-84 represent antisense oligonucleotides directed against CC transforming growth factor beta-1 (TGF beta-1). The oligonucleotides CC exemplify the invention. The specification describes oligonucleotides CC that contain 8-30 nucleotides, which contain at most 8 nucleotides that CC can each form three hydrogen bonds to cytosine; do not contain four CC consecutive nucleotides able to form three H-bonds each to four CC consecutive cytosines; do not contain two sequences of three consecutive cytosines; do not contain two sequences of three consecutive cytosines, and the ratio between residues able to form three H-bonds to three consecutive CC cytosines, and the ratio between residues able to form two H-bonds each CC (2R) or three such bonds (3R) is given by 2R/JR = 0.33-0.72. The CC cytosines, are used to modulate expression of genes, particularly CC the genes for p53, ErbB-2, junB, junD, TGF-beta 1 or beta 2 to control CC proliferation of primary cell cultures (e.g. bone marrow stem, liver or continue cells, osteoclaste, osteoblasts and/or keratinocytes). The CC altering their expression or activity) and therapeutically, e.g. in cases of first contain cases for p53, Farberson or activity) and therapeutically, e.g. in cases
                                                                                                             Human genome, biallelic marker; high density disequilibrium map, genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation; haplotyping; hybridisation; identification; SNP; PCR primer;
                                                                                                                                                                                                                                                               Human biallelic marker downstream amplification primer SEQ ID NO:11752
                                                                                                                                                                                                                                                                                                                                            10-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ77396
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Preparation of antisense oligo:nucleotide(s) which lack long runs of consecutive guanosine or inosine - and have specific ratio of residues able to form two or three hydrogen bonds, have greater activity and reduced toxicity, used therapeutically or to modulate growth of cells:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1;
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Homo sapiens

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Best Local S
Matches 13
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                                                                                                                     Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; bolymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation; haplotyping; hybridisation; dentification; SNP; PCR primer; amplification; single nucleotide polymorphism; SNP; PCR primer;
                                                                                                                                                                                                                                     AAZ75225 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleotide sequences. AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-APR-1998;
23-NOV-1998;
                                                                                                            diagnosis; ss.
                                                                                                                                                                         Human
                                                                                                                                                                                            10-SEP-2001
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 21-APR-1998;
23-NOV-1998;
                                                                       WO9954500-A2
                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies
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                              21-APR-1999;
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 98US-0082614P.
98US-0109732P.
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98US-0109732P.
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                              99WO-IB000822
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                                                                                                                                                                                                                                                                                                                                                                     C; 7
                                                                                                                                                                       downstream amplification
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Pred. No. 4.2e+04;
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RESULT 62
AAF83133/c
ID AAF83133
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Best Local S
Matches 13
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       Chandrasekhar
                                              (ELIL )
(MILE/)
(ONYI/)
(THIR/)
                                                                                                                                                                                                       26-SEP-2000; 2000WO-US026497
                                                                                                                                                                                                                                                    05-APR-2001.
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                                                                                                                                                                                                                                                                                                                                             Synthetic
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                                            ) LILLY & CO ELI.
) MILES R R.
) ONYIA J E.
) THIRUNAVUKKARASU
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    Halladay
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100.0%; Pr
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Pred. No.
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4.2e+04;
hes 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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Best Local
         Isolated nucleic acid fragment comprising the transcriptional regulatory region of the human osteoprotegerin (opg) gene, useful for identifying compounds which are useful in the treatment of bone disease, arthritis
                                                                                                                                                                                                                                                                                                                                                                             Osteoprotegerin; opg; human; transcriptional regulation; bone disease; arterial disease, immune function; lymph node development; T-cell; B-cell; antiarthritic; immunomodulatory; antirheumatic; antiinflammatory;
                                                                         WPI; 2001-281682/29
                                                                                                        Chandrasekhar
                                                                                                                                                 (ELIL)
                                                                                                                                                                                                                27-SEP-1999;
                                                                                                                                                                                                                                             26-SEP-2000; 2000WO-US026497
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                                                                                                                                                                                                                                                                                                                                        Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence of primer 17897469 S6.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Isolated nucleic acid fragment comprising the transcriptional regulatory region of the human osteoprotegerin (opg) gene, useful for identifying compounds which are useful in the treatment of bone disease, arthritis
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                                                                                                                                 ) LILLY & CO ELI.
') MILES R R.
') ONYIA J E.
') THIRUNAVUKKARASU K.
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                                                                                                                                                                                                                                                                                                                                                                   primer;
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                                                                                                    Halladay
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Pred. No.
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                                                                                                      Martin
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. 4.2e+04;
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RESULT 64
AASO2440/c
ID AASO244
XX AASO244
XX AASO244
XX AASO24
XX Human
XX Human
XX Human
XX AASO24
XX AASO24
DT 18-JUL
DE Human
XX Human
XX AASO24
DT 27-SEP
DR W02001
PR 27-JUN
PP 27-SEP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention provides the transcriptional regulatory region of the human osteoprotegerin (opg) gene, and fragments exhibiting human opg gene transcriptional regulatory activity. The isolated opg nucleic acid fragments are useful in an assay to identify an agonist or antagonist of opg expression. They are also useful for the manufacture of a composition for the diagnosis of a human susceptible to, predisposed to, or at concern of the diagnosis of of a composition of or increased risk for developing a symptom, condition, or disease caused by cover- or under-expression of opg. The agonist, antagonist and modulators are useful in the treatment of a disease in a human caused by abnormal expression of opg. The disease in a human caused by abnormal concession of opg. The disease is bone disease, arthritis, arterial concession of opg. The disease is bone disease, arthritis, arterial concess, abnormal immune function, abnormal lymph node development, or sequences ARRS3129-136 represent PCR primers which are not related to the present invention. The specification has certain pages missing and the above mentioned primers are a part of pages from other patent
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Best Local S
Matches 13
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27-JUN-2000;
26-SEP-2000;
                                                                                   Example 1;
                                                                                                                                                                                                                                                                                                              Shimkets
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   angiogenesis; immune associated disorder; gestational disopre-eclampsia; neuronal development; immunogen; antibody; agonist; TSR1; sequencing primer; 17897469 S5; ss.
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                                                                                                                                                              TSRX PRO (PRO comprising thror identify molecules modulating
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                                                                                                                                                                                                                                                    2001-266157/27.
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                                                                                                                                                                                                                                                                                                                                                                  CURAGEN CORP
                                                                                                                                       rheumatoid
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                                                                                Page
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 3 A; 6 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                      2000US-0214759P
2000US-00669360
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                                                                                                                                                                                                                                                                                                           Vernet
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000US-0214759P.
                                                                                89; 116pp;
                                                                                                                                          arthritis
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                                                                                                                                                                 thrombospondin-I repeat) domain useful ating TSRX activity or function, for tro
                                                                                                                                                                                                                                                                                                        Tchernev
                                                                                English
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Pred. No.
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4.2e+04;
                                                                                                                                       neovascularization
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The sequence represents a sequencing primer used clone encoding the Human thrombospondin-1 repeat

to sequence the cDNA (TSR) domain contain

containing

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RESULT 65
AAS02441
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Best Local S
Matches 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; Thrombospondin repeat domain; TSRX; cancer; breast cancer; rheumatoid arthritis; ocular neovascularisation; wound healing; angiogenesis; immune associated disorder; gestational disorder; pre-eclampsia; neuronal development; immunogen; antibody; antisense; agonist; TSR1; sequencing primer; 17897469 S6; ss.
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27-JUN-2000; 2000US-0214759P.
26-SEP-2000; 2000US-00669360.
                                                        TSRX PRO (PRO comprising thrombospondin-I repeat) domain useful to identify molecules modulating TSRX activity or function, for treat: cancer, rheumatoid arthritis and ocular neovascularization.
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                                                                                                                                                                                WPI; 2001-266157/27.
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                                                                                                                                                                                                                                                                                           (CURA-) CURAGEN CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequencing primer 17897469
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                                                                                                                                                                                                                                   Vernet
     , 68
116pp;
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                                                                                                                                                                                                                                   Tchernev
  English.
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Pred. No.
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                                                                                                                                                                                                                                   Boldog
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                                                                                                                                                                                                                                   Herrmann
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The sequence represents a sequencing primer used to sequence the cDNA clone encoding the Human thrombospondin-1 repeat (TSR) domain contain

domain containing

antisense therapy

The present invention describes the human Zmax1 gene and the high bone mass (HBM) gene, which are found on chromosome 11q13.3. The Zmax1 and H genes have osteopathic activities. The genes can be used in gene thera

and in the production of

vaccines.

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used therapy Disclosure; Page 36; 443pp;

English

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RESULT 66
ABA82453
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Best Local (
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                                                                                                                                               New high bone mass (HBM) modulating bone mass for
                                                                                                                                                                                                                                                                                                                                             05-APR-2000;
05-APR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200177327-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Zmax1
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                                                                                                                                                                                                                                                                                                     (GENO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       are used for tissue typing and forensic identification
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene region physical map preparation STS marker #412
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13; Conserv
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                                                                                                                                                                                                                                                                                                     GENOME THERAPEUTICS CORP.
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Conservative
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                                                                                                                                                                                                                                                            Little RD,
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2000US-00544398.
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                                                                                                                                                                                                                                                          Recker RR,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 13;
Pred. No.
                                                                                                                                                 Zmaxl genes and proteins useful treatment of e.g. osteoporosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            porosis; osteopathic; gene therapy;
disorder; Paget's disease; adapter;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zmax1 gene;
                                                                                                                                                                                                                                                            Johnson
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4.2e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           chromosome 11;
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diagnosis and treatment of bone disorders including osteoporosis,

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KESULT 67
ABK23250
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The invention relates to a method for identifying a molecule involved in CC lipid regulation comprising identifying a molecule that binds to or inhibits binding of a molecule to high bone mass (HBM) or its wild type gene, Zmaxl. Compounds identified by the method are useful for treating, CC disgnosing, preventing or screening for normal and abnormal lipid-associated conditions, including arteriosclerosis, cardiovascular disease, stroke, and osteoporosis. The compounds may also be used in the treatment or prevention of diabetic atherosclerosis, neurovascular conditions caused by plaque build-up, poor circulation due to plaque build-up and associated poor wound healing. The methods may be used in gene thorapy, pharmaceutical development, and diagnostic assays for bone development disorders. Molecules identified by comparison of Zmaxl and HBM systems can be used as surrogate markers in pharmaceutical conditions can be used as surrogate markers in pharmaceutical conditions can be used as surrogate markers in pharmaceutical conditions can be used as surrogate markers, probes, linkers treatment of bone diseases. Sequences ABK22776-ABK23411 represent cDNA molecules encoding human Zmaxl and HBM, and PCR primers, probes, linkers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; mouse; Zmax1; HBM; high bone mass gene; lipid regulation; stroke; lipid-associated condition; arteriosclerosis; cardiovascular disease; ss; cardiovascular disease; cardiovascular disease; ss; cardiovascular disease; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Identifying molecules involved in lipid regulation, useful for diagnosing, treating or preventing e.g., arteriosclerosis, comidentifying a molecule that binds to high bone mass gene or it corresponding wild type gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   osteoporosis; atherosclerosis; diabetic atherosclerosis; plaque build-up neurovascular condition; wound healing; gene therapy; PCR primer; probe; bone development disorder; antiarteriosclerotic; cardiovascular;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Paget's disease, sclerostosis, osteomalacia and fibrous dysplasia. ABA82038 to ABA82700 and AAG68168 to AAG68193 represent sequences used the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 41; 409pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Carulli JP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-APR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR primer
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Pred. No.
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4.2e+04;
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RESULT 68
ACC45833
      The invention relates to novel transgenic animals expressing the high CC bone mass (HBM) gene, expressing the corresponding wild type HBM gene, CC comprising an alteration of the gene encoding LRP5 or LRP6, or expressing CC an LRP5 that is modulated by an altered gene control sequence introduced CC by homologous or non-homologous recombination. The transgenic animals are CC for the study of bone density modulation or bone mass modulation. The cinvention has osteopathic and cytostatic activity. The polynucleotides of CC invention may have a use in gene therapy. The transgenic animals and CC mucleic acids are for the study of bone density modulation, where the CC bone mass is modulated relative to non-transgenic animals of the same CC species in more than one parameter selected from bone density, bone CC transgenic animals, nucleic acids and methods are useful for identifying CC molecules involved in bone development, and for developing pharmaceutical compositions, which may be employed for treating or preventing bone CC neoplasms of the bone. The transgenic animals and nucleic acids are also cases, in hone development.
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17-MAY-2001; 2001US-0291311P.
01-FEB-2002; 2002US-0353058P.
04-MAR-2002; 2002US-0361293P.
                                                                                                                                                                                                                                                                                                                                                    New transgenic animals (e.g. mice), useful as models density modulation, developing drugs for treating or diseases (e.g. osteoporosis), or diagnosing diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Babij P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; high bone mass; HBM; LRP5; LRP6; transgenic; bone mass modulation; gene therapy; bone density modulation; bone strength; trabecular number; bone size; bone tissue connectivity; bone disease; osteoprossis; PCR; osteomalacia; rickets; Paget's disease; neoplasm of the bone; primer; ss.
                                                                                                                                                                                                                                                                                                         Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACC45833 standard;
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      methods
                                                                                                                                                                                                                                                                                                       Page 57;
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                                                                                                                                                                                                                                                                                                                                          density.
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 for diagnosing
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Pred. No.
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development,
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RESULT 69
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                                                             Query Match
Best Local S
Matches 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
                                                                                                                                                                      The present invention relates to High Bone Mass (HBM), LRP5 (Zmax1) and LRP6 mutants, which results in a HBM-like phenotype when expressed in a cell. The HBM-like phenotype results in bone mass modulation and/or lipid level modulation. The invention is useful for diagnosing a HBM-like phenotype in a subject and for preparing a composition for modulating bone mass and/or lipid levels in a subject suffering from e.g. osteoporosis. The present sequence is a Sequence Tagged Site (STS) marker, which was used to prepare a physical map of the Zmax1 (LRP5) gene
                                                                                                                                                                                                                                                                                                                                                          New nucleic acid comprising a mutation diagnosing a HBM-like phenotype in a su composition for modulating bone mass ar suffering from e.g. osteoporosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-MAY-2001; 2001US-0290071P.
17-MAY-2001; 2001US-0291311P.
01-FEB-2002; 2002US-0353058P.
04-MAR-2002; 2002US-0361293P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  or characterised by used in the exemplif
                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                             Example 2; Page 64; 629pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-129214/12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Allen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GENO-) GENOME THERAPEUTICS CORP
(AMHP) WYETH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                pathic; Gene therapy;
mass modulation; oster
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 203
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                                                               l Similarity
13; Conserv
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                         ACAAGTGTCTGGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Anisowicz
                                                                                                                                BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; 2001US-0291311P.
; 2002US-0353058P.
; 2002US-0361293P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.6%;
ilarity 100.0%;
Conservative
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                                                             larity 100.0%;
Conservative (
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                                                                                                                             4 C; 6
                                                          2.6%; Score 13; DB
100.0%; Pred. No. 4.;
ive 0; Mismatches
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osteoporosis; STS; sequence tagg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Graham JR,
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                                                                                                                           G; 3 T; 0 U; 0 Other;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                              a subject
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Morales
                                                                                                                                                                                                                                                                                                                                                                           on in LRP5 or LRP6, useful for subject and for preparing a and/or lipid levels in a subject
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                                                             4.2e+04;
hes 0;
                                                                                                DB 10;
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hes 0;
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                                                                                            Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Yaworsky
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                                                             Indels
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ged site; ds.
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                                                             Gaps
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8
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                                                                                                                                                                                            RESULT 71
ABK11325
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ABZ75622
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Best Local S
Matches 13
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                                                                                                             05-JUN-2002
                                                                                                                                              ABK11325;
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                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19
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involves using a multiplex polymerase chain reaction assay having chromosome-specific short tandem repeat (STR) markers. The STR marker 21 328 (informal designation) is useful as a marker for the diagnosis of aneuploidy of a chromosome, particularly trisomy 21, 13, 18 or X, or Turner Syndrome. The STR marker Y-40S (informal designation) is useful a marker for the diagnosis of the sex of an individual. Marker CF508 is useful for detecting the presence or absence of a genetic disease, particularly cystic fibrosis. Sequences ABZ75621-22 represent PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Detecting aneuploidy of a chromosome in a fetus by using a multiplex polymerase chain reaction assay comprising chromosome-specific short tandem repeat markers.
ss; PCR; ACH4; Acyl coenzyme A thioesterase; plant; transgenic;
lipid oxidation; regulation of Coenzyme A; fatty acid metabolism; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Aneuploidy; chromosome; multiplex assay; polymerase chain reaction; short tandem repeat; STR; turner syndrome; cystic fibrosis; primer;
                                                                      Arabidopsis Acyl coenzyme A thioesterase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to detecting aneuploidy of a chromosome and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 16; 30pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-FEB-2001; 2001GB-00004690
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STR marker 21-32S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-MAY-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                     standard;
                                                                                                                                                                                                                                                                                                                                            CCTGGGTGACAAG 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry
                                                                                                                      (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STR marker 21-32S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      specific
                                                                                                                                                                                                                   DNA;
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                                                                                                                        entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      6 C; 6 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCR
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 6; Le
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                                                                        PCR primer ACH4-LBamHI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to an isolated acyl coenzyme A thioesterase (ACH) encoding nucleic acid, encoding one of ACH1, ACH2, ACH4 or ACH5. ACH enzymes have a role in lipid oxidation, regulation of Coenzyme A pools and in fatty acid metabolism. Also include are a host cell transfected with the nucleic acid, a transgenic plant transfected with the nucleic acid (including its seed or oil) and ACH antisense molecules. The ACH nucleic acid is useful for altering a phenotype of a plant and for making a transgenic plant, by transfecting the plant tissue with the ACH nucleic acid under conditions such that a transgenic plant is generated. The ACH nucleic acid is also useful for producing variants of acyl-CoA thiosterace.
                                                                                                                                                    PCR; primer; ss;
DNA high-density
microarray; Type
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel acyl coenzyme A thioesterase gene useful for altering a phenotype of a plant, making a transgenic plant and for producing variants of acyl-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-JUL-2000; 2000US-0220028P
16-JUL-2001; 2001US-00906408
              05-OCT-2001; 2001US-00972469
                                        05-OCT-2001; 2001US-00972469
                                                                                                US2003073085-A1
                                                                                                                            Homo sapiens
                                                                                                                                                                                                             Sense PCR primer used to amplify CDC37.
                                                                                                                                                                                                                                        16-OCT-2003 (first entry)
                                                                                                                                                                                                                                                                                             ACH66467 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19 BP; 5 A; 4 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     thioesterases. The present sequence is a PCR primer used to amplify Arabidopsis ACH4 encoding sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Page 47; 78pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CoA thioesterases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Arabidopsis thaliana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (TILT/) TILTON G B. (SHOC/) SHOCKEY J M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (BROW/) BROWSE J A.
                                                                                                                                                                                                                                                                                                                                                                                             250 ATCCTGAAGAATT 262
                                                                                                                                                                                                                                                                                                                                                                                                                     1 Similarity
13; Conserv
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                                                                                                                                                                                                                                                                                                                                                                 N
                                                                                                                                                                                                                                                                                                                                                                 ATCCTGAAGAATT 14
                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shockey JM,
                                                                                                                                                    genomic DNA; gDNA; untranslated region; UTR; microarray; biosite; large scale production; II primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                    2.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Browse
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Pred. No
                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                     No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 6;
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                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                   gDNA probe;
                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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cc species. The method involves identifying a 3' untranslated region (UTR) cc of a gDNA sequence, designing probe, performing PCR, separating the cc product by size differentiation and performing PCR, separating the cc product by size differentiation and performing a second PCR to amplify cc the predetermined sequence. Also claimed is a biological analysis device, cc gequences, located on the substrate, which are generated by the method cc above and a DNA high-density microarray comprising a substrate upon which are generated by the method cc are deposited an array of biosites of genomic DNA fragments having the sequence, where the genomic DNA fragments have a sequence length of from cc sequences, where the genomic DNA fragments have a sequence length of from capturing in microarray formats, fabricates high-density DNA arrays of large quantities of gDNA probes, which enables greater efficiency for cc printing in microarray formats, fabricates high-density DNA arrays of derived sequences by simple PCR amplifications without cloning. The method produces amplified sequences that have greater specificity and cc size consistency than that observed with cDNA fragments, and allows for greater signal sensitivity than oligonucleotides. The sequence presented cc is a Type II gene specific primer
The invention discloses a method for amplifying expressed genetic sequences from genomic DNA (GDNA) from mammalian or higher order plant sequences from genomic DNA (GDNA) from mammalian or higher order plant sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Amplifying expressed genetic sequences from genomic DNA of mammalian or higher order plant species for printing on DNA microarrays, involves using the 3' untranslated region of the gene sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 7; 15pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lai F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (LAIF/) LAI F. (ZHOU/) ZHOU D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Zhou D;
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Query Match Best Local S Matches 13 Similarity Conservative 100.0%; 2.6%; Score 13; DB; Pred. No. 4.2 0, DB 9; 4.2e+04; 0 Length 19

0

Sequence 19

BP; 2 A; 8 C; 3 G; 6 T; 0 U; 0 Other;

밁 Ş 36 CTGCTTCCAGCCC 48 μ. CTGCTTCCAGCCC 13 Indels 0, Gaps

RESULT 73 ADO14975 standard; RNA; 19 BP

Human PDGFr-targeted siNA lower strand SEQ ID NO:406

01-JUL-2004

(first entry)

ADO14975/c
ID ADO149
XX
AC ADO149
XX
O1-JUL
XX
Human
XX
Cytost
KW treati
KW bronch
KW short
KW short
KW expres
KW cytost
KW expres
KW platel
XX
PN WO2003
XX
PD 04-SEF
XX short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; treating leukaemia; solid tumors; restenosis; polycystic kidney disease; bronchiolitis; glomerulonephritis; stroke; RNA interference; cytostatic; vasotropic; platelet derived function analysis; growth gene mapping; human factor receptor; PDO nephrotropic; cerebroprotective; PDGFr; ss

WO2003072704-A2

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                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local
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06-JUN-2002;
29-AUG-2002;
cytostatic; vasotropic; nephrotropic; cerebroprotective; treating leukaemia; solid tumors; restenosis; polycystic kidney disease; bronchiolitis; glomerulonephritis; stroke; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
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09-SEP-2002;
                                                                                                                                                                                                                                    ADO14664 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to short interfering nucleic acids (siMA) which downregulate expression of the human platelet-derived growth factor receptor (PDGFr) gene by RNA interference. The siMAS may or may not comprise ribonucleotides and may be double or single stranded. They
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                                                                                                                 Human PDGFr-targeted siNA upper strand SEQ ID NO:95.
                                                                                                                                                         01-JUL-2004
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2002US-0386782P.

2002US-0406784P.

2002US-0406784P.

2002US-0408378P.

2002US-0409293P.

2003US-0440129P.
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Pred. No. 4.2e+04;
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RESULT 75 ADM82819/c ID ADM828 XX

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standard; DNA; 19

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Matches Query Match Best Local

10;

Conservative

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Mismatches

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Gaps

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Similarity

2.6%;

Score 13; Pred. No. 4.2e+04;

DB 11; Length 19

108

GGTCTGCCTGCAG 120

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CC downregulate expression of the human platelet-derived growth factor comprise ribonucleotides and may be double or single stranded. They comprise ribonucleotides and may be double or single stranded. They comprise ribonucleotides and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the sinAs include short interfaring RNA (siRNA, double-cc stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain cc deoxyribonucleotides, and can be chemically synthesised, expressed from a cyettor or enzymatically synthesised. The invention also relates to kits cromplexes of siRNA; and vectors that express siNA. The siNAs are used to complexes of siRNA; and vectors that express siNA, ribe siNAs are used to corganisms (e.g., by ex vivo gene therapy), or in grafts and transplants cordisates, bronchiolitis, glomerulonephritis and stroke. The siNAs are also useful for drug screening, diagnosis, therapputic target identification and validation, genetic engineering, pharmacogenomics, studying gene c function, and gene mapping (e.g., of single nucleotide polymorphisms).

The present sequence represents the upper strand of a human pDGFr transcript cargeted double-stranded siNA, which is identical to the PDGFr transcript
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05-SEP-2002;
09-SEP-2002;
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11-MAR-2002;
06-JUN-2002;
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therapeutic target identification; pharmacog
gene function analysis, gene mapping, human,
platelet derived growth factor receptor; PDC
Sequence 19
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pharmacogenomics;
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ADM82819;

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CC although known, have not previously been associated with schizophrenia, CC although known, have not previously been associated with schizophrenia, CC stress responses (which can precipitate exacerbate the above conditions). CC The genes can be used for screening an individual with schizophrenia, CC ADHD and/or bipolar disorder, for monitoring the progression or treatment of these conditions, for identifying agents useful in the treatment of continuous conditions, for identifying agents useful in the treatment of continuous conditions, for identifying agents useful in the treatment of continuous conditions, for identifying agents useful in the treatment of continuous conditions, for identifying agents useful in the treatment of continuous conditions and for treating or continuous conditions and for treatment of continuous conditions and continuous conditions and continuous conditions and for treatment of continuous conditions continuous conditions and for the invention continuous cont
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Use of known genes associated with schizophrenia, attention-deficit-hyperactivity disorder and bipolar disorder for screening, monitoring progression or treating schizophrenia, attention-deficit-hyperactivity disorder or bipolar disorder.
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22-MAY-2003; 2003US-0472489P
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NOVARTIS PHARMA GMBH.
UNIV MARYLAND BALTIMORE.
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Sequence 19

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Best Local Similarity
Matches 13; Conserv
19 GCCCATGCCCGAC
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4.2e+04;
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                                      Sequence 7840, Ap

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Sequence 37, Appl

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Sequence 26, Appli
Sequence 29, Appl
Sequence 19, Appl
Sequence 192, Appl
Sequence 193, Appl
Sequence 193, Appl
Sequence 10366, A
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RESULT 1 US-10-422-523-21

ALIGNMENTS

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Sequence 21, Application US/10422523

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APPLICANT: Latif, Farida
APPLICANT: Wei, Ming-Hui
APPLICANT: Wei, Ming-Hui
APPLICANT: Sekido, Yoshitaka
APPLICANT: Geo, Boning
APPLICANT: Duh, Fuh-Mei
ITILE OF INVENTION: Calcium Channel Compositions and Methods of Use T
FILE REFERENCE: NIH-05043
CURRENT APPLICATION NUMBER: US/10/116,949
CURRENT APPLICATION NUMBER: APPLICATION NUMBER: US/09/470,443
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US/09/470,443
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/114,359
PRIOR FILING DATE: EARLIER FILING DATE: 1998-12-30
NUMBER OF SEQ ID NOS: 114
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 37
LENGTH: 22
RESULT 3
US-10-215-112-6487
; Sequence 6487, Application US/10215112
; Publication No. US20030082596A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                 ; FEATURE: ; OTHER INFORMATION: Description of Artificial Sequence: US-10-116-949-37
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US-10-116-949-37
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APPLICANT: SHORT, JAY M.
APPLICANT: SHORT, JAY M.
TITLE OF INVENTION: SYNTHETIC LIGATION REASSEMBLY IN DIRECTED EVOLUTION
FILE REFERENCE: DIV-1460-15A US
CURRENT PAPLICATION NUMBER: US/10/422,523
CURRENT FILING DATE: 2003-04-24
PRIOR APPLICATION NUMBER: 09/332,835
PRIOR APPLICATION NUMBER: 09/332,835
PRIOR FILING DATE: 1999-06-14
NUMBER OF SEQ ID NOS: 36
SOFTWARE: PATENTIN VET. 2.1
SEQ ID NO 21
LENGTH: 21
TYPEE. DATE
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Best Local Similarity
Matches 15; Conserv
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APPLICANT: Minna, John D.
APPLICANT: Latif, Farida
APPLICANT: wei, Ming-Hui
APPLICANT: Sekido, Yoshitaka
                                                                                                                                                                                                                                                                                                                                         LENGTH: 22
TYPE: DNA
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
FEATURE:
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IO. US20030044911A1
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1.7e+03;
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US-10-261-517-12/c
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US-10-215-112-6487
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FILE REFERENCE: 13425-032001
CURRENT APPLICATION NUMBER: US/10/261,517
CURRENT FILING DATE: 2002-10-01
PRIOR APPLICATION NUMBER: US/09/645,629
PRIOR FILING DATE: 2000-08-24
PRIOR APPLICATION NUMBER: US 60/151,867
PRIOR APPLICATION NUMBER: US 60/151,867
PRIOR APPLICATION NUMBER: SE 9903009-0
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PRIOR FILING DATE: 2001-03-16
NUMBER OF SEQ ID NOS: 131066
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 20798
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20798, Application US/10098263B Publication No. US20030104410A1 GENERAL INFORMATION:
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CURRENT FILLING DATE: 2002-08-08
NUMBER OF SEQ ID NOS: 14936
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 6487
LENGTH: 25
                                                                                                                                                                                                                                                                                              Sequence 12, Application US/10261517 Publication No. US20030143583A1
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CURRENT APPLICATION NUMBER: US/10/098,263B
CURRENT FILING DATE: 2003-01-08
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                                                                                                                                                                                                            APPLICANT: Climent-Johansson, Isabel
APPLICANT: Dahlman-Wright, Karin
APPLICANT: Lake, Staffan
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                                                                                                                                                                 APPLICANT: Wasserman, Wyeth
TITLE OF INVENTION: NOVEL RESPONSE ELEMENT
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TYPE: DNA
ORGANISM: Homo sapien
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ORGANISM: Artificial Sequence
FEATURE:
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larity 100.0%; Pred. No. 1.7e+0;
Conservative 0; Mismatches
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100.0%; Pred. No.
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; TYPE: RNA
; ORGANISM: Mus musculus
US-10-440-850-367
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US-10-440-850-367
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OTHER INFORMATION: synthetically generated oligonucleotide
US-10-261-517-12
                                                                                                                                                                                                                                                                             RESULT 7
US-10-712-672-1858/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/440,850
CURRENT FILING DATE: 2003-05-19
PRIOR APPLICATION NUMBER: US/09/650,012
PRIOR FILING DATE: 2000-08-28
PRIOR APPLICATION NUMBER: US 08/585,684
PRIOR APPLICATION NUMBER: US 08/585,684
PRIOR APPLICATION NUMBER: US 60/000,951
PRIOR APPLICATION NUMBER: US 60/000,951
PRIOR FILING DATE: 1995-07-07
PRIOR APPLICATION NUMBER: US 09/038,073
PRIOR APPLICATION NUMBER: US 09/038,073
PRIOR FILING DATE: 1998-03-11
NUMBER OF SEQ ID NOS: 2285
SOFTWARE: Patentin version 3.0
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LENGTH: 25
                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                 Sequence 1858, Application US/10712672 Publication No. US20040102413A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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Best Local Similarity 78.6%;
Matches 11; Conservative
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Best Local Similarity
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                            APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
TITLE OF INVENTION: Method and Reagent for the Inhibition of Telomerase Enzyme
FILE REFERENCE: MBHB00-882-C (400/019)
CURRENT APPLICATION NUMBER: US/10/712,672
CURRENT FILING DATE: 2003-11-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Method and Reagent TITLE OF INVENTION: Immune Responses FILE REFERENCE: 250/110 (MBHB00-900-A)
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                                                                                                                                                                   APPLICANT: Ribozyme Pharmaceuticals, Inc APPLICANT: Chowrira, Bharat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Ribozyme Pharmaceuticals, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
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PRIOR FILING DATE: 2000-08-31
                      PRIOR APPLICATION NUMBER: US/09/653,225
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b. US20030207837A1
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Pred. No.
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CURRENT APPLICATION NUMBER: US/09/891.517
CURRENT FILING DATE: 2001-06-27
PRIOR APPLICATION NUMBER: UP2000-193133
PRIOR FILING DATE: 2000-06-27
PRIOR APPLICATION NUMBER: JP2000-236115
PRIOR FILING DATE: 2000-08-03
PRIOR APPLICATION NUMBER: JP2000-292483
PRIOR APPLICATION NUMBER: JP2000-292483
PRIOR FILING DATE: 2000-99-26
NUMBER OF SEQ ID NOS: 108
SOFTWARE: PATENTIN VEYSION 3.1
SEQ ID NO 99
                                                                   US-10-751-736-36439/c
; Sequence 36439, Application US/10751736
; Publication No. US20040265230A1
; GENERAL INFORMATION:
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; OTHER INFORMATION: Synthetic DNA
US-09-891-517-99
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US-10-712-672-1858
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Best Local Similarity
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Patent No. US20020106653A1
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APPLICANT:
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PRIOR TILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/150,713
PRIOR FILING DATE: 1999-08-31
NUMBER OF SEQ ID NOS: 5586
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: KURATA, SHINYA
APPLICANT: YAMADA, KAZUTAKA
APPLICANT: YOKOMAKU, TOYOKAKU
TITLE OF INVENTION: NOVEL NUCLEIC ACID PROBES,
TITLE OF INVENTION: NUCLEIC ACID BY USING THE
TITLE OF INVENTION: METHOD
 APPLICANT:
                                  APPLICANT: Wyeth APPLICANT: Mart
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: KURANE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: 210352US-1994-163-0-X
                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA ORGANISM: Artificial Sequence
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Brown, En
Liu, Wei
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KAMAGATA, YOICHI
TORIMURA, MASAKI
                                    Martinez, Robert
                                                                                                                                                                                                                                                                                 Conservative
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                 Eugene
                                                                                                                                                                                                                                                                                             2.8%; Score 14;
100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                              Length 19;
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Sequence 36640, Application US/10751736

Publication No. US20040265230A1

GENERAL INFORMATION:
APPLICANT: Wyeth
APPLICANT: Brown, Eugene
APPLICANT: Brown, Eugene
APPLICANT: Liu, Wei
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TRE
TITLE OF INVENTION: CANCERS
FILE REFERENCE: AM100927 (031896-002000)
CURRENT APPLICATION NUMBER: US/10/751,736
CURRENT FILING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US Provisional Application 60/438,000
PRIOR FILING DATE: 2003-01-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 54873
SOFTWARE: Patentin version 3.2
SEQ ID NO 36440
LENGTH: 21
TYPE: RNA
OPCANTON
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; ORGANISM: RNAi
US-10-751-736-36440
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APPLICANT: Wyeth
APPLICANT: Martinez
APPLICANT: Brown, E
APPLICANT: Liu, Wei
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US-10-751-736-36440/c
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CURRENT APPLICATION NUMBER: US/10/751,736
CURRENT FILING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US Provisional Application 60/438,000
PRIOR FILING DATE: 2003-01-06
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100.0%; Pred. No. 6.:
ive 0; Mismatches
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hes 0;
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                                                                                                                         FOR DIAGNOSING AND TREATING COLON
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US-10-751-736-36641/c
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US-10-751-736-47110
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; ORGANISM: RNA1
US-10-751-736-36641
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SOFTWARE: Patentin version 3.2
SEQ ID NO 36640
LENGTH: 21
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SEQ ID NO 36641
LENGTH: 21
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Best Local Similarity 100.0%;
Matches 14; Conservative (
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SEQ ID NO 47110
LENGTH: 21
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Best Local :
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APPLICANT: Liu, Wei
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING COLON
TITLE OF INVENTION: CANCERS
FILE REFERENCE: AN100927 (031896-002000)
CURRENT APPLICATION NUMBER: US/10/751,736
CURRENT FILING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US Provisional Application 60/438,000
PRIOR APPLICATION NUMBER: US PROVISIONAL APPLICATION NUMBER: US PROVIS
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING COLON
TITLE OF INVENTION: CANCERS
FILE REFERENCE: AM10927 (031896-002000)
CURRENT APPLICATION NUMBER: US/10/751,736
CURRENT FILING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US Provisional Application 60/438,000
PRIOR PLING DATE: 2003-01-06
NUMBER OF SEQ ID NOS: 54873
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TYPE: DNA
ORGANISM: homo sapiens
10-751-736-47110
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; LENGTH: 21
; TYPE: RNA
; ORGANISM: RNAi
US-10-751-736-47111
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APPLICANT: Martinez, Robert

APPLICANT: Brown, Eugene

APPLICANT: Liu, Wei

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TRE

TITLE OF INVENTION: CANCERS

FILE REFERENCE: AM100927 (031896-002000).

CURRENT APPLICATION NUMBER: US/10/751,736

CURRENT FILING DATE: 2003-01-06

PRIOR APPLICATION NUMBER: US Provisional Application 60/438,000

PRIOR FILING DATE: 2003-01-06

NUMBER OF SEQ ID NOS: 54873

SOFTWARE: PatentIn version 3.2
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                                                                                                                                               SOFTWARE: PatentIn version 3.2
SEQ ID NO 47113
LENGTH: 21
TYPE: DNA
ORGANISM: homo sapiens
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Best Local Similarity 100.
Matches 14; Conservative
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Best Local Similarity 78.6%;
Matches 11; Conservative
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                                                                         Query Match
Best Local &
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APPLICANT: Martinez, Robert
APPLICANT: Brown, Eugene
APPLICANT: Liu, Wei
TITLE OF INVENTION: COMPOSITIONS AND METHODS
TITLE OF INVENTION: CANCERS
FILE REFERENCE: AM100927 (031896-002000)
                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: US Provisional Application 60/438,000 PRIOR FILING DATE: 2003-01-06 NUMBER OF SEQ ID NOS: 54873
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CURRENT FILING DATE: 2003-01-06
395 GCAGCTTCCGTGAG 408
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                                                                           Similarity
                                                           Conservative
                                                                         2.8%; Score 14;
100.0%; Pred. No.
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100.0%; Pred. No.
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Pred. No. 6.1e+03;
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6.1e+03;
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RESULT 18
US-10-142-662-3
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US-10-751-736-47114
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; ORGANISM: RNAi
US-10-751-736-47114
Sequence 3, Application US/10142662
Publication No. US20030211462A1
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Publication No. US20030211462A1
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Best Local Similarity
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APPLICANT: Yu, Simon
TITLE OF INVENTION: DIRECTIONALLY CLONED RANDOM CDNA EXPRESSION VECTOR LIBRARIES,
TITLE OF INVENTION: COMPOSITIONS AND METHODS OF USE
TILE REFERENCE: A-71241/RMS/DHR
CURRENT FILLORION NUMBER: US/10/142,662
CURRENT FILLING DATE: 2002-12-10
NUMBER OF SEQ ID NOS: 53
SOFTWARE: Patentin version 3.1
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APPLICANT: Liu, Wei
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING COLON
TITLE OF INVENTION: CANCERS
FILE REFERENCE: AM100927 (031896-002000)
CURRENT APPLICATION NUMBER: US/10/751,736
CURRENT FILING DATE: 2003-01-06
FRIOR APPLICATION NUMBER: US Provisional Application 60/438,000
RIOR FILING DATE: 2003-01-06
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APPLICANT: Wu, Xian
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                                                                                                                                                                                                                                                                                                                TYPE: DNA
ORGANISM: Artificial sequence
FEATURE:
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Pred. No. 6.
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US-10-325-694-19/c
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                                                                                                                                                                                                                                                                                                                                                                                       Best Local
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; OTHER INFORMATION: synthetic US-10-142-662-3
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APPLICANT: Wu, Xian
APPLICANT: Wu, Xian
APPLICANT: Shen, Mary
APPLICANT: Yu, Simon
TITLE OF INVENTION: DIRECTIONALLY CLONED RANDOM CDNA EXPRESSION VECTOR LIBRARIES,
TITLE OF INVENTION: COMPOSITIONS AND METHODS OF USE
FILE REFERENCE: A-71241/RMS/DHR
CURRENT APPLICATION NUMBER: US/10/142,662
CURRENT FILING DATE: 2002-12-10
NUMBER OF SEQ ID NOS: 53
SOFTWARE: Patentin version 3.1
SEQ ID NO 3
                                                                                                                                                                           Sequence 4828, Application US/10032585 Publication No. US20030180953A1 GENERAL INFORMATION:
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APPLICANT: Terry, Roemer D.
APPLICANT: Bo, Jiang
APPLICANT: Charles, Boone
APPLICANT: Charles, Boone
TITLE OF INVENTION: Gene Disruption Methodologies
FILE REFERENCE: 10182-005-999
CURRENT APPLICATION NUMBER: US/10/032,585
CURRENT FILING DATE: 2001-12-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 19, Application US/10325694 Publication No. US20030148463A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 24
TYPE: DNA
ORGANISM: HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: KUFER, PETER
APPLICANT: RAUM, TOBLE
TITLE OF INVENTION: NOVEL METHOD FOR THE PRODUCTION OF ANTI-HUMAN ANTIGEN
TITLE OF INVENTION: RECEPTORS AND USES THEREOF
FILE REFERENCE: 38164000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US/09/403,107
PRIOR FILING DATE: 1999-10-14
NUMBER OF SEQ ID NOS: 152
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CURRENT FILING DATE: 2002-12-19
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TYPE: DNA
ORGANISM: Artificial sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                         185 GTGAGCTCAACCTG 198
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Similarity 100.0%;
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100.0%; Pred. No.
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Pred. No.
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0. 6.1e+03;
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. 6.1e+03;
                                                                    for Drug Target Discovery
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 24;
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Sequence 12370, Application US/10215112

Publication No. US20030082596A1

GENERAL INFORMATION:
APPLICANT: Michael Mittmann
ITITLE OF INVENTION: Method of Genetic Analysis of Probes:
TITLE OF INVENTION: Test3

FILE REFERENCE: 3119

CURRENT APPLICATION NUMBER: US/10/215,112

CURRENT FILING DATE: 2002-08-08

NUMBER OF SEQ ID NOS: 14936

SOFTWARE: FastSEQ for Windows Version 4.0

LENGTH: 25
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US-10-098-263B-125035/c
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                                                                                                                            ; TYPE: DNA
; ORGANISM: Homo sapien
US-10-098-263B-125035
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US-10-215-112-12370
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US-10-215-112-12370
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Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                               SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 125035
LENGTH: 25
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LENGTH: 24
                                                        Matches
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Best Local Similarity
                                                                                       Query Match
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TITLE OF INVENTION: Human Microarray
FILE REFERENCE: 31.8.1
CURRENT APPLICATION NUMBER: US/10/098,263B
CURRENT FILING DATE: 2003-01-08
PRIOR APPLICATION NUMBER: 60/276,759
PRIOR FILING DATE: 2001-03-16
NUMBER OF SEQ ID NOS: 131066
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SOFTWARE: PatentIn version 3.1
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ORGANISM: Candida albicans
                                                    Local Similarity
les 14; Conserv
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53 CCGACCGGAGAGCG 66
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                                                    2.8%; Score 14; DB 15; Length 25; 100.0%; Pred. No. 6.1e+03; tive 0; Mismatches 0; Indels
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100.0%; Pred. No. 6.1e+03
tive 0; Mismatches 0
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RESULT 25 US-10-112-917-6

GENERAL INFORMATION:

APPLICANT: Adams, Se

Sean H

TITLE OF INVENTION: FIZZ1 FOR METABOLISM REGULATION FILE REFERENCE: 9800080-0002

Sequence 6, Application US/10112917 Publication No. US20030104351A1

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; LENGTH: 25
; TYPE: DNA
; ORGANISM: probe
US-10-775-169-1156
                                                                                                                                                                                                                           ; APPLICANT: Burczynski, Michael
; APPLICANT: Twine, Natalie
; APPLICANT: Twine, Natalie
; APPLICANT: Trepicchio, William
; TITLE OF INVENTION: Method for Monitoring Dr
; FILE REFERENCE: AM101800 (031896-013000)
; CURRENT APPLICATION NUMBER: US/10/775,169
; CURRENT FILING DATE: 2004-02-11
; NUMBER OF SEQ ID NOS: 5278
; SOFTWARE: Patentin version 3.2
; SEQ ID NO 1156
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US-10-775-169-1156
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US-10-775-169-1155
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APPLICANT: Dorner, Andrew
APPLICANT: Trepicchio, William
APPLICANT: Trepicchio, William
TITLE OF INVENTION: Method for Monitoring Drug Activities In Vivo
FILE REFERENCE: AM101080 (031896-013000)
CURRENT APPLICATION NUMBER: US/10/775,169
CURRENT FILING DATE: 2004-02-11
NUMBER OF SEQ ID NOS: 5278
SOFTWARE: PatentIn version 3.2
SEQ ID NO 1155
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Publication No. US20040175743A1
                                                                          Matches
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                                                                                                              Query Match
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Wyeth
                                                                                          / Match 2.8%; Score 14; DB 17; Local Similarity 100.0%; Pred. No. 6.1e+03
                      94 CCTCAGCAGTGTGC 107
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                                                                          14; Conservative
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                                                                          Mismatches
                                                                                            6.1e+03;
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                                                                                                            Length 25;
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                                                                                                                                                                                                                                                                                  RESULT 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; OTHER INFORMATION: synthetic US-10-142-662-5
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       GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
APPLICANT: HANZEL, David K.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
CURRENT FILING DATE: 2001-05-25
                                                                                                                                                                                                         Sequence 7839, Application US/09866108
Patent No. US20020048800A1
GENERAL INFORMATION:
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LENGTH: 27
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LENGTH: 26
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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Publication No. US20030211462A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: YL, SIMON
TITLE OF INVENTION: DIRECTIONALLY CLONED RANDOM CDNA
TITLE OF INVENTION: COMPOSITIONS AND METHODS OF USE
FILE REFERENCE: A-71241/RMS/DHR
CURRENT APPLICATION NUMBER: US/10/142,662
CURRENT FILING DATE: 2002-12-10
NUMBER OF SEQ ID NOS: 53
NUMBER OF SEQ ID NOS: 53
SOFTWARE: Patentin version 3.1
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APPLICANT: Wu, Xian
APPLICANT: Shen, Maz
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PRIOR FILING DATE: 2001-03-30
NUMBER OF SEQ ID NOS: 10
SOFTWARE: PatentIn version 3.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/10/112,917
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA ORGANISM: Artificial sequence
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OTHER INFORMATION: oligonucleotide probe
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                                                                                                                                                                                                                                                                                                                                                                                                                            Local
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APPLICATION NUMBER: US 60/207,456
                                                                                                                                                                                                                                                                                                                                                                       450 GCCAGTGGCCGTAA 463
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                                                                                                                                                                                                                                                                                                                                    GCCAGTGGCCGTAA 6
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                                                                                                                                                                                                                                                                                                                                                                                                                        2.8%;
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Pred. No.
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                               FILE REFERENCE: AECMICA-7

CURRENT APPLICATION NUMBER: US/09/866,108

CURRENT APPLICATION NUMBER: US/09/866,108

CURRENT APPLICATION NUMBER: US 60/207,456

PRIOR FILLING DATE: 2000-05-26

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR FILLING DATE: 2000-09-27

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR APPLICATION NUMBER: PCT/US01/00666

PRIOR APPLICATION NUMBER: PCT/US01/00667

PRIOR APPLICATION NUMBER: PCT/US01/00667
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GENERAL INFORMATION:
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Best Local :
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SEQ ID NO 7839
                                                                                                                                                                                                                                                                                         APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
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OR APPLICATION NUMBER: GB 24263.6
OR FILING DATE: 2000-10-04
OR APPLICATION NUMBER: US 60/236,359
OR FILING DATE: 2000-09-27
OR APPLICATION NUMBER: PCT/US01/00666
OR APPLICATION NUMBER: PCT/US01/00667
OR FILING DATE: 2001-01-30
OR FILING DATE: 2001-01-30
OR FILING DATE: 2001-01-30
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Gaps

DATE: 2001-01-30

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RESULT 29
US-09-866-108-7841/c
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; ORGANISM: Homo sapiens
US-09-866-108-7840
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PRIOR FILING DATE: 2000-05-26
PRIOR PPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR PILING DATE: 2000-10-04
PRIOR PPLICATION NUMBER: US 60/236,359
PRIOR PILING DATE: 2000-09-20
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00664
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APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharro
APPLICANT: HANZEL, David
APPLICANT: RANK, David
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Best Local Similarity
Matches 13; Conserv
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                                                                                                     PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/866,108
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APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
FILE REFERENCE: AEOMICA-7
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                                                              APPLICATION NUMBER: PCT/US01/00668 FILING DATE: 2001-01-30
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APPLICATION NUMBER: US 60/266,860
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APPLICATION NUMBER: PCT/US01/00668
FILING DATE: 2001-01-30
                         APPLICATION NUMBER: PCT/US01/00663 FILING DATE: 2001-01-30
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FILING DATE:
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ICATION NUMBER: PCT/US01/00670
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RANK, David R.
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RESULT 30
US-09-866-108-7842/c
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PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR PILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
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PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR PRIOR DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 2000-09-21
PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
NUMBER OF SEQ ID NOS: 15752
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Patent No.
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SEQ ID NO 7841
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Best Local Similarity 100.0%;
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CURRENT FILING DATE: 2001-05-25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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ORGANISM: Homo sapiens
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
FILING DATE: 2001-01-30
                   FILING DATE: 2000-09-21
APPLICATION NUMBER: US 60/266,860
FILING DATE: 2001-02-05
                                                                                                 APPLICATION NUMBER: PCT/US01/00661
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00670
FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00662
FILING DATE: 2001-01-30
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SEQ ID NOS: 15752
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NN, Sharron
NZEL, David
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Pred. No.
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; ORGANISM: Homo sapiens
US-09-866-108-7843
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; ORGANISM: Homo sapiens
US-09-866-108-7842
Query Match
Best Local Similarity
Matches 13; Conserv
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LENGTH: 17
                                                                                                                                         SOFTWARE: Aeomica
SEQ ID NO 7843
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Best Local Similarity
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PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: ABOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
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                                                                                                                                                                                           PRIOR FILING DATE:
                                                                                                     TYPE: DNA
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ITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
ILE REFERENCE: AEOMICA-7
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FILING DATE: 2001-01-30
FILING DATE: 2001-01-30
                                                                                                                                                                                                           FILING DATE: 2000-09-21
APPLICATION NUMBER: US 60/266,860
                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: PCT/US01/00661 FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00669
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JI, Yonggang
PENN, Sharron G.
PENN David K.
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     Conservative
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Score 13; DB 9; Le; Pred. No. 2.2e+04; 0; Mismatches 0;
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Pred. No.
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APPLICANT: SHANNON, MARK
TITLE OF INVENTION: HUMAN MYOSIN-LIKE POLYPEPTIDE EXPRESSED PREDOMINANTLY IN HEART AN
FILE REFERENCE: PB0105
CURRENT APPLICATION NUMBER: US/10/723,361
CURRENT APPLICATION NUMBER: US/03-11-26
PRIOR PILING DATE: 2003-01-26
PRIOR FILING DATE: 2001-05-26
PRIOR FILING DATE: 2001-05-26
PRIOR PILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
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; Sequence 677, App.
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APPLICANT: Ribozyme Pharmaceuticals,
APPLICANT: McSwiggen, James
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Best Local Similarity
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APPLICANT: JI
APPLICANT: PE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Publication No.
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APPLICANT:
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TITLE OF INVENTION: Nucleic Acid Treatment
FILE REFERENCE: 400/057 (MBHB01-1158-A)
CURRENT APPLICATION NUMBER: 02 10/238,700
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ORGANISM: Homo sapiens
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APPLICATION NUMBER: PCT/US01/00665
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PENN, Sharron
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; ORGANISM: Homo sapiens
US-10-723-361-7840
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; ORGANISM: Homo sapiens
US-10-723-361-7839
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PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR APPLICATION NUMBER: PCT/US01/00665
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: PB0105

CURRENT APPLICATION NUMBER: US/10/723,361

CURRENT FILING DATE: 2003-11-26

PRIOR APPLICATION NUMBER: US 09/866,108

PRIOR PILING DATE: 2001-05-25

PRIOR APPLICATION NUMBER: US 60/207,456

PRIOR FILING DATE: 2000-05-26

PRIOR APPLICATION NUMBER: GB 24263.6

PRIOR APPLICATION NUMBER: GB 0/236,359

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR APPLICATION NUMBER: PCT/US01/00666
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NUMBER OF SEQ ID NOS: 15755
SOFTWARE: Acomica Sequence Listing Engine
SEQ ID NO 7839
LENGTH: 17
                                                                                    Query Match
Best Local
                                                                                                                                                                                                                SEQ ID NO 7840
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SOFTWARE: Acomica Sequence Listing Engine
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TITLE OF INVENTION: HUMAN MYOSIN-LIKE POLYPEPTIDE EXPRESSED PREDOMINANTLY IN HEART
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                                 CACCIGCIGCITC 42
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                                                                      Conservative
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                                                                                    Score 13; pred. No.
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Pred. No.
                                                                    Mismatches
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                                                                                  DB 17; Length 17; 2.2e+04;
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2.2e+04;
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US-10-723-361-7842/c
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Best Local :
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                                 APPLICANT: SHANNON, MAIK
TITLE OF INVENTION: HUMAN MYOSIN-LIKE POLYPEPTIDE
FILE REFERENCE: PBO105
CURRENT APPLICATION NUMBER: US/10/723,361
CURRENT FILING DATE: 2003-11-26
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PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
                                                                                                                                                                 APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggan
APPLICANT: PENN, Shar:
APPLICANT: HANZEL, Dav
PRIOR APPLICATION NUMBER: US 09/866,108 PRIOR FILING DATE: 2001-05-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Aeomica Sequence Listing Engine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Remaining Prior Application data removed - NUMBER OF SEQ ID NOS: 15755
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CURRENT FILING DATE: 2003-11-26
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APPLICANT: SHANNON, MARK
TITLE OF INVENTION: HUMAN MYOSIN-LIKE POLYPEPTIDE EXPRESSED PREDOMINANTLY IN HEART 1
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APPLICANT: JI, Yonggang
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TYPE: DNA
ORGANISM: Homo sapiens
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FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00669
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APPLICATION NUMBER: PCT/US01/00667
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ion No. US20040137589A1
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PENN, Sharron G
HANZEL, David K
RANK, David R.
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HANZEL, David
RANK, David R.
                                                                                                                            CHEN, Wensheng
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APPLICATION NUMBER:

2000-05-26

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GB 24263.6

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US-10-723-361-7842
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CURRENT FILING DATE: 2003-11-26
PRIOR APPLICATION NUMBER: US 09/866,108
PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
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Best Local
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                   PRIOR APPLICATION NUMBER: PCT/US01/00668
                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: SHANNON, MAIK
TITLE OF INVENTION: HUMAN MYOSIN-LIKE POLYPEPTIDE EXPRESSED PREDOMINANTLY IN HEART
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PRIOR APPLICATION NUMBER: PCT/US01/00669
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00668
FILING DATE: 2001-01-30
                                                                                                                                                                         APPLICATION NUMBER: PCT/US01/00666
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00667
FILING DATE: 2001-01-30
                                                                           APPLICATION NUMBER: PCT/US01/00665
                                                                                             APPLICATION NUMBER: PCT/US01/00669 FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00664
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                                                          FILING DATE: 2001-01-30
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EL, David
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APPLICANT: Cohen, Daniel
APPLICANT: Chumakov, Ilya
APPLICANT: Chumakov, Ilya
APPLICANT: Chumakov, Ilya
TITLE OF INVENTION: Biallelic markers for use in construc
FILE REFERENCE: GENSET, 020CP1
CURRENT APPLICATION NUMBER: US/10/349,143
CURRENT FILING DATE: 2003-01-21
PRIOR APPLICATION NUMBER: US/09/422,978
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US
PRIOR FILING DATE: EARLIER APPLICATION NUMBER: US
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PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US
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US-90-349-143-9581/c
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US-10-349-143-11752/c
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US-10-723-361-7843
                                                                                                                                                                                                                                                  Sequence 11752, Application US/10349143
Publication No. US20040005584A1
GENERAL INFORMATION:
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LENGTH: 17
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PRIOR PILING DATE: 1999-10-20
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER:
PRIOR FILING DATE: EARLIER FILING DATE: 1999-04-21
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER:
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER:
PRIOR FILING DATE: EARLIER FILING DATE: 1998-11-23
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER:
PRIOR FILING DATE: EARLIER FILING DATE: 1998-04-21
PRIOR FILING DATE: EARLIER FILING DATE: 1998-04-21
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APPLICANT: Blumer
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CURRENT FILING DATE: 2003-01-21
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LOCATION: 1..18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
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Pred. No.
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Pred. No.
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2.2e+04;
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                                                                                                                                                                     in constructing a high density...
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          US 60/109,732
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RESULT 41
US-10-477-238A-478
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US-10-731-739-478
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                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                    Sequence 478, Application US/10477238A Publication No. US20040221326A1
                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 478
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PRIOR APPLICATION NUMBER: GARLIER APPLICATION NUMBER: US 60/082,614
PRIOR FILING DATE: EARLIER FILING DATE: 1998-04-21
NUMBER OF SEQ ID NOS: 11796
SEQ ID NO 11752
ENERTH: 18
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Best Local :
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APPLICANT: Little, Randall D.
APPLICANT: Recker, Robert R.
APPLICANT: Johnson, Mark L.
TITLE OF INVENTION: High bone mass gene of
FILE REFERENCE: 032796-013
CURRENT APPLICATION NUMBER: US/10/731,739
CURRENT FILING DATE: 203-12-10
               APPLICANT: Yaworsky, Paul
APPLICANT: Bex, Frederick J. III
APPLICANT: Bodine, Peter Van Nest
TITLE OF INVENTION: Transgenic Animal Model of Bone Mass Modulation
FILE REFERENCE: 032796-212
                                                       APPLICANT: Babij, Philip
APPLICANT: Yaworsky, Pa
APPLICANT: Bex, Frederi
APPLICANT: Bodine, Pete
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CURRENT APPLICATION NUMBER: US/10/477,238A
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NAME/KEY: primer_bind
LOCATION: 1..18
OTHER INFORMATION: downstream amplification primer 99-5075 for SEQ
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100.0%; Pred. No.
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RESULT 43

US-09-906-408A-37

Sequence 37, Application US/09906408A Publication No. US20030028915A1 GENERAL INFORMATION:

APPLICANT: Tilton,

Gregory

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CURRENT FILING DATE: 2003-10-08

PRIOR APPLICATION NUMBER: PCT/USO2/14876

PRIOR APPLICATION NUMBER: US 60/290,071

PRIOR APPLICATION NUMBER: US 60/290,071

PRIOR FILING DATE: 2001-05-11

PRIOR APPLICATION NUMBER: US 60/291,311

PRIOR APPLICATION NUMBER: US 60/291,311

PRIOR APPLICATION NUMBER: US 60/353,058

PRIOR APPLICATION NUMBER: US 60/353,058

PRIOR APPLICATION NUMBER: US 60/361,293

PRIOR FILING DATE: 2002-03-04

NUMBER OF SEQ ID NOS: 812

SOFTMARE: PastSEQ for Windows Version 4.0

SEQ ID NO 478

LENGTH: 18
                                                                                                                                                                                    ; ORGANISM: Homo sapiens US-10-680-287A-478
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PRIOR FILING DATE: 2001-05-11
PRIOR PELICATION NUMBER: US 60/291,311
PRIOR FILING DATE: 2001-05-17
PRIOR FILING DATE: 2001-05-17
PRIOR APPLICATION NUMBER: US 60/353,058
PRIOR FILING DATE: 2002-02-01
PRIOR APPLICATION NUMBER: US 60/361,293
PRIOR FILING DATE: 2002-03-04
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Best Local Similarity
Matches 13; Conserv
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 478
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: 032796-179
CURRENT APPLICATION NUMBER: US/10/680,287A
CURRENT FILING DATE: 2003-10-08
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                                                                                                                                                                                                                                 TYPE: DNA
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PPLICANT: Bex, Frederick J. III
PPLICANT: Bodine, Peter Van Nest
ITLE OF INVENTION: Transgenic Animal Model of Bone Mass Modulation
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                                          203 ACAAGTGTCTGGA 215
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                                                                                       13; Conservative 0;
2 ACAAGTGTCTGGA 14
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o. US20040244069A1
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100.0%; Pred. No.
ative 0; Mismatc)
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100.0%; Pred. No. 2.
ive 0; Mismatches
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RESULT 45
US-10-224-005-101/c
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; ORGANISM: Homo sapiens
US-09-972-469-61
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US-09-972-469-61
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PRIOR FILLING DATE: 2000-07-21
NUMBER OF SEQ ID NOS: 46
SOFTWARE: Patentin version 3.0
SEQ ID NO 37
LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 13; Conserv
                                                                                                                                                                                                                                              Sequence 101, Application US/10224005 Publication No. US20030143732A1 GENERAL INFORMATION:
SEQ ID NO 101
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                                           APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: Fosnaugh, Kathy
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Adenosine Al Receptor
TITLE OF INVENTION: Gene Expression Using Short Interfering RNA
TILE REFERENCE: 900/041 (MBHB01-1110-A)
CURRENT APPLICATION NUMBER: US/10/224,005
CURRENT FILING DATE: 2002-08-20
PRIOR APPLICATION NUMBER: US 60/315,315
PRIOR PILING DATE: 2001-08-28
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                                 NUMBER OF SEQ ID NOS: 347
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CURRENT FILING DATE: 2001-10-05
NUMBER OF SEQ ID NOS: 196
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APPLICANT: Zhou, Dai
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CURRENT APPLICATION NUMBER: US/09/906,408A
CURRENT FILING DATE: 2001-07-16
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OTHER INFORMATION: Synthetic
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ORGANISM: Artificial Sequence
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                 PatentIn version 3.0
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FILE REFERENCE: 900/041 (MBHB01-1110-A)

FULE REFERENCE: 900/041 (MBHB01-1110-A)

CURRENT APPLICATION NUMBER: US/10/224,005

CURRENT FILING DATE: 2002-08-20

PRIOR APPLICATION NUMBER: US 60/315,315

PRIOR TLING DATE: 2001-08-28

NUMBER OF SEQ ID NOS: 347

SOPTWARE: Patentin version 3.0

SEQ ID NO 262

1.ENCORT.
                                           TITLE OF INVENTION: GENE INVOLVED IN CADASIL, ME
TITLE OF INVENTION: THERAPEUTIC APPLICATION
FILE REFERENCE: 03715.0048-00000
CURRENT APPLICATION NUMBER: US/10/356,625
CURRENT FILING DATE: 2003-02-03
PRIOR APPLICATION NUMBER: US/09/230,652
PRIOR APPLICATION NUMBER: US/09/230,652
PRIOR APPLICATION NUMBER: FR 96 09733
PRIOR FILING DATE: 1996-08-01
PRIOR FILING DATE: 1996-08-01
PRIOR FILING DATE: 1997-04-16
PRIOR APPLICATION NUMBER: FR 97 04680
PRIOR APPLICATION NUMBER: FR 97 04680
PRIOR FILING DATE: 1997-04-16
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                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                            Sequence 100, A Publication No.
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Best Local (
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              PRIOR APPLICATION NUMBER: PCT/FR97/01433 PRIOR FILING DATE: 1997-07-31
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                                                                                                                                                                                                                                                                                                                                APPLICANT: Tournier-Lasserve, Elisabeth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: McSwiggen, James APPLICANT: Fosnaugh, Kathy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: RNA Interference Mediated Inhibition of Adenosine Al Receptor TITLE OF INVENTION: Gene Expression Using Short Interfering RNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
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OF SEQ ID NOS: 163
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Bousser, Marie-Germaine
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o. US20030186290A1
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Pred. No. 2.2e+04
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Pred. No.
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2.2e+04;
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RESULT 49
US-10-160-787-97
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US-10-160-787-20/c
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                                                                                                                                  ; FEATURE:
US-10-160-787-97
                                                                                                                                                                                 SEQ ID NO 97
LENGTH: 20
TYPE: DNA
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APPLICANT: ANDREW T. MALE
TITLE OF INVENTION: ANTISENSE MODULATION OF PCTAIRE PROTEIN KINASE 2 EXPRESSION
FILE REFERENCE: RTS-0204
                                                                Query Match
Best Local Similarity
Matches 13; Conserv
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Best Local
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CURRENT FILING DATE: 2002-05-31
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CURRENT FILING DATE: 2002-05-31
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TITLE OF INVENTION: ANTISENSE MODULATION OF PCTAIRE PROTEIN KINASE 2 EXPRESSION
FILE REFERENCE: RTS-0204
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SEQ ID NO 100
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SEQ ID NO 20
LENGTH: 20
                                                                                                                                                          ORGANISM: H. sapiens
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CATGTTGACCGAG 15
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                                                              DB 15; Length 20;
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ches 0; Indels
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APPLICANT: C. Frank Bennett
APPLICANT: Kenneth W. Dobie
TITLE OF INVENTION: MODULATION OF PHOSPHODIESTERASE 8A
FILE REFERENCE: PTS-0062
CURRENT APPLICATION NUMBER: US/10/318,389
CURRENT FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 134
SEQ ID NO 23
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
EEATURE:
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US-10-318-389-94/c
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APPLICANT: C. Frank Bennett
APPLICANT: Kenneth W. Dobis
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SEQ ID NO 94
LENGTH: 20
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                                                                                                                                                                              Sequence 94, Application US/10318389
Publication No. US20040121328A1
                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local :
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Best Local S
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APPLICANT: Kenneth W. Dobie
TITLE OF INVENTION: ANTISENSE MODULATION OF
FILE REFERENCE: RTS-0365
CURRENT APPLICATION NUMBER: US/10/159,856
CURRENT FILING DATE: 2002-05-31
NUMBER OF SEQ ID NOS: 134
                                  CURRENT APPLICATION NUMBER: US/10/318,389
CURRENT FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 134
                                                                                     APPLICANT: Kenneth W. Dobie
TITLE OF INVENTION: MODULATION OF PHOSPHODIESTERASE
FILE REFERENCE: PTS-0062
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TYPE: DNA
ORGANIEM: Artificial Sequence
FEATURE:
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Similarity 100.0%;
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Similarity 100.0%; Pred. No. 2.2e+04
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Pred. No. 2.2e+04
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; TYPE: RNA
; ORGANISM: RNAi
US-10-751-736-3929
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US-10-751-736-3929
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; ORGANISM: Homo sapiens
US-10-184-085A-763
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US-10-184-085A-763/c
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                                                                                                                                                               TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING COLON
TITLE OF INVENTION: CANCERS
FILE REFERENCE: AM100927 (031896-002000)
CURRENT APPLICATION NUMBER: US/10/751,736
CURRENT FILING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US Provisional Application 60/438,000
PRIOR PRIOR DATE: 2003-01-06
NUMBER OF SEQ ID NOS: 54873
SOFTWARE: PatentIn version 3.2
SEQ ID NO 3929
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Matches
Query Match
Best Local Similarity
Matches 9; Conserv
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Publication No. US20040265230A1
GENERAL INFORMATION:
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SEQ ID NO 763
LENGTH: 21
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APPLICANT: Marti
APPLICANT: Brown
APPLICANT: Liu,
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APPLICANT: Luebke, Kevin, J.
APPLICANT: Balog, Robert P.
TITLE OF INVENTION: Identification of Chemically Modified Polymers
FILE REFERENCE: 119999-1035
CURRENT APPLICATION NUMBER: US/10/184,085A
CURRENT FILING DATE: 2002-10-01
PRIOR APPLICATION NUMBER: US 60/301,370
PRIOR FILING DATE: 2001-06-27
NUMBER OF SEQ ID NOS: 1291
COURTENEDS SECOND NUMBER: US 60/301,370
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                                                                                                                                                   LENGTH: 21
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Local Similarity 100.0%; Pred. No. 2.2e+04;
heB 13; Conservative 0; Mismatches 0; Indels
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Liu, Wei
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o. US20030152950A1
    Conservative
                       2.6%;
                   Score 13; DB 18;
Pred. No. 2.2e+04;
    Mismatches
                                        DB 18; Length 21;
  Indels
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; ORGANISM: homo sapiens
US-10-751-736-11683
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US-10-751-736-11684
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US-10-751-736-11683
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING COLON
TITLE OF INVENTION: CANCERS
FILE REFERENCE: AM100927 (031896-002000)
CURRENT APPLICATION NUMBER: US/10/751,736
CURRENT FILING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US Provisional Application 60/438,000
PRIOR FILING DATE: 2003-01-06
PRIOR FILING DATE: 2003-01-06
NUMBER OF SEQ ID NOS: 54873
SOFTWARE: PAtentin version 3.2
SEQ ID NO 11684
LENGTH: 21
TYPE: RNA.
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FILE REFERENCE, AM100927 (031896-002000)
CURRENT APPLICATION NUMBER: US/10/751,736
CURRENT FILING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US Provisional Application 60/438,000
PRIOR FILING DATE: 2003-01-06
PRIOR FILING DATE: 2003-01-06
OVER PRIOR OF SEQ ID NOS: 54873
SOFTWARE: Patentin version 3.2
SEQ ID NO 11683
LENGTH: 21
                                                                     Query Match 2.6%;
Best Local Similarity 69.2%;
Matches 9; Conservative
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Best Local Similarity
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APPLICANT: Marti
APPLICANT: Brown
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Pred. No.
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RESULT 57 US-10-751-736-11686

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US-10-751-736-36637/c
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US-10-751-736-11687
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US-10-751-736-11687
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; ORGANISM: homo sapiens
US-10-751-736-11686
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CURRENT APPLICATION NUMBER: US/10/751,736
CURRENT FILING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US PROVISIONAL Application 60/438,000
PRIOR PILING DATE: 2003-01-06
NUMBER OF SEQ ID NOS: 54873
SOFTWARE: Patentin version 3.2
SEQ ID NO 11686
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SEQ ID NO 11687
LENGTH: 21
                                                                         Sequence 36637, Application US/10751736 Publication No. US20040265230A1 GENERAL INFORMATION:
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APPLICANT: Marti
APPLICANT: Brown
APPLICANT: Liu,
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Publication No. US20040265230A1
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FILE REFERENCE: AM100927 (031896-002000)
CURRENT APPLICATION NUMBER: US/10/751,736
CURRENT FILING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US Provisional Application 60/438,000
PRIOR FILING DATE: 2003-01-06
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NUMBER OF SEQ ID NOS: 54873
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Local Similarity 69.2%;
tes 9; Conservative
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Local Similarity 100.0%; Pred. No.
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                                                           Wyeth
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Martinez, Robert
Brown, Eugene
Liu, Wei
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Pred. No. 2.2e+04;
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; ORGANISM: homo sapiens
US-10-751-736-36637
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CURRENT APPLICATION NUMBER: US/10/751,736

CURRENT FILING DATE: 2003-01-06

PRIOR APPLICATION NUMBER: US Provisional Application 60/438,000

PRIOR FILING DATE: 2003-01-06

NUMBER OF SEQ ID NOS: 54873

SOPTWARE: Patentin version 3.2

SEQ ID NO 36637
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SOFTWARE: PatentIn version 3.2
SEQ ID NO 36638
LENGTH: 21
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APPLICANT: Liu, Wei
ITTLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING COLON
TITLE OF INVENTION: CANCERS
FILE REFERENCE: AM10927 (031896-002000)
CURRENT FILING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US Provisional Application 60/438,000
PRIOR PILING DATE: 2003-01-06
APPLICANT: Liu, Wei
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND
TITLE OF INVENTION: CANCERS
FILE REFERENCE: AM100927 (031896-002000)
CURRENT APPLICATION NUMBER: US/10/751,736
CURRENT FILING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US Provisional Application 60/438,000
PRIOR FILING DATE: 2003-01-06
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2.2e+04;
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                                                                                                                                        DIAGNOSING AND TREATING COLON
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; TYPE: DNA
; ORGANISM: homo sapiens
US-10-751-736-49795
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CURRENT APPLICATION NUMBER: US/10/751,736
PRIOR APPLICATION NUMBER: US/10/751,736
PRIOR FILING DATE: 2003-01-06
PRIOR FILING DATE: 2003-01-06
PRIOR FILING DATE: 2003-01-06
NUMBER OF SEQ ID NOS: 54873
SOFTWARE: Patentin version 3.2
SEQ ID NO 49746
LENGTH: 21
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US-10-751-736-49795/c
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US-10-751-736-49746
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; ORGANISM: homo sapiens
US-10-751-736-36643
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                                                                     SOFTWARE: PatentIn version 3.2 SEQ ID NO 49795
                                                                                                                                                                                                                                                                                                                    Sequence 49795, Application US/10751736 Publication No. US20040265230A1 GENERAL INFORMATION:
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SOFTWARE: PatentIn version 3.2
SEQ ID NO 36643
LENGTH: 21
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                                                                                                                                                   APPLICANT: Brown, Eugene
APPLICANT: Liu, Wei
TITLE OF INVENTION: COMPOSITIONS AND METHODS
TITLE OF INVENTION: CANCERS
FILE REFERENCE: AM100927 (031896-002000)
CURRENT APPLICATION NUMBER: US/10/751,736
CURRENT FILING DATE: 2003-01-06
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING COLON TITLE OF INVENTION: CANCERS
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TYPE: RN
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ORGANISM: RNAi
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Liu, Wei
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61.5%;
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Pred. No. 2.2e+04;
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2.2e+04;
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US-10-751-736-49798
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US-10-751-736-49796
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US-10-751-736-49796/c
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                                                                                                                                              SEQ ID NO 49798
LENGTH: 21
TYPE: DNA
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SOFTWARE: PatentIn version 3.2
SEQ ID NO 49796
LENGTH: 21
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                                                   Matches
                                                                               Query Match
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Best Local Similarity
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APPLICANT: Marti
APPLICANT: Brown
APPLICANT: Liu,
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Best Local 9
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SOFTWARE: PatentIn version 1 2
EQ ID NO 40700
                                                                                                                                                                                                                       TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING COLON TITLE OF INVENTION: CANCERS FILE REFERENCE: AM100927 (031896-002000) CURRENT APPLICATION NUMBER: US/10/751,736 CURRENT FILING DATE: 2003-01-06 PRIOR APPLICATION NUMBER: US Provisional Application 60/438,000 PRIOR FILING DATE: 2003-01-06
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                                                                                                                                                                                                                                                                                                                                      APPLICANT: Brown, Eugene
APPLICANT: Liu, Wei
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                                                                  Local
28 TGCACCTGCTGCT 40
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                               2.6%; Score 13;
Similarity 100.0%; Pred. No.
13; Conservative 0; Microco
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100.0%; Pred. No.
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                                                             DB 18;
0. 2.2e+04;
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2.2e+04;
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TGCACCTGCTGCT 2

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US-10-203-939-46
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; ORGANISM: Melon
US-10-203-939-46
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SEQ ID NO 46
LENGTH: 22
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Publication No.
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TITLE OF INVENTION: Nucleic Acid Sequences and Differrent Uses Thereof
FILE REFERENCE: BAR ILAN UNIV. (Chen)
CURRENT APPLICATION NUMBER: US/10/203,939
CURRENT FILING DATE: 2002-11-19
CURRENT APPLICATION NUMBER: US/10/236,392
CURRENT FILING DATE: 2002-09-06
PRIOR APPLICATION NUMBER: US09/540,763
PRIOR FILING DATE: 2000-03-30
                                                                                  APPLICANT: Reiger, Daniel K
APPLICANT: Reiger, Daniel K
APPLICANT: Rothenberg, Mark E
APPLICANT: Shenberg, Mark E
APPLICANT: Shenby, Suresh
APPLICANT: Shimkets, Richard A
APPLICANT: Shimkets, Richard A
APPLICANT: Smithson, Glennda
APPLICANT: Smithson, Glennda
APPLICANT: THERAPEUTIC POLYPEPTIDES, NUCLEIC ACIDS ENCODING SAME
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                                                                   FILE REFERENCE: 21402-442A
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Malyankar, Uriel M
Miller, Charles E
Millet, Isabelle
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Gusev, Vladamir
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Crabtree, Julie
Edinger, Shlomit, R
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Vo. US20030172396A1
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o. US20040067490A1
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William M
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; TYPE: DNA
; ORGANISM: Artificial Sequence
; PEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Forward Primer
US-10-236-392-297
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                                                                                                                       US-10-806-782-5
                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US/09/587,945
PRIOR FILLING DATE: 2000-06-06
PRIOR APPLICATION NUMBER: US 60/190,692
PRIOR FILLING DATE: 2000-03-20
PRIOR FILLING DATE: 1998-05-26
PRIOR FILLING DATE: 1998-05-26
PRIOR APPLICATION NUMBER: SE 9701963-2
PRIOR FILLING DATE: 1997-05-26
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LENGTH: 22
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Best Local Similarity 100.0%;
Marches 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 5, Application US/10806782 Publication No. US20040166061A1 GENERAL INFORMATION:
                                                          Query Match
Best Local (
                                         Matches
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CURRENT FILING DATE: 2004-03-23
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NUMBER OF SEQ ID NOS: 794
SOFTWARE: Custom
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE: 10806-117A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Carlsson, Peter TITLE OF INVENTION: Animal Model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Enerback, Sven
APPLICANT: Carlsson, Pet
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                                                                                                                                                                     TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                       OTHER INFORMATION: Description
                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                         LENGTH: 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US60/318,120 FILING DATE: 2001-09-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US60/369,479 FILING DATE: 2002-04-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US60/390,155 FILING DATE: 2002-06-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US60/318,130
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US60/357,303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US60/318,765
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  115 CTGCAGCCTTTCT 127
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                                                            Similarity
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                                       Conservative
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                                   2.6%; Score 13;
100.0%; Pred. No.
tive 0; Mismatch
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                                                                                                                                     of Artificial Sequence: 5' primer
                                       Mismatches
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                                                          DB 17; I
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2.2e+04;
                                                                             Length 22;
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APPLICANT: HUH, Tae-Lin

APPLICANT: PARK, Hae-Chul

APPLICANT: KIM, Chul-Hee

APPLICANT: KIM, Hyung-seok

TITLE OF INVENTION: STRUCTURAL GENES, TRANSGENIC ANIMAL HAVING HUC PROMOTER AND ITS

TITLE OF INVENTION: METHOD FOR SCREENING NEURONAL MUTANT ANIMALS USING THE TRANSGENI

CURRENT APPLICATION NUMBER: US/10/296,665

CURRENT FILING DATE: 2002-11-25

NUMBER OF SEQ ID NOS: 3

SOFTWARE: Patentin version 3.1

IENGTH: 24

TYPE: """
                                                                                                           ; FEATURE:
; NAME/KEY: primer bind
; LOCATION: (1)..(74)
; OTHER INFORMATION:
US-10-296-665-2
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                                                 Matches
                                                                               Query Match
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SEQ ID NO 12
LENGTH: 23
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Publication No. US20030236194A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: CAI, Jidong
APPLICANT: ARDATI, Mohamad Ali
APPLICANT: SANDRASAGRA, Anthony
TITLS OF INVENTION: A NOVEL G PROTEIN-COUPLED RECEPTOR, GAVE 2
FILE REFERENCE: USAV2001/0051 USNP
CURRENT APPLICATION NUMBER: US/10/438,657
CURRENT FILING DATE: 2003-05-15
PRIOR APPLICATION NUMBER: US 60/382,375
PRIOR FILING DATE: 2002-05-23
NUMBER: OF SEQ ID NOS: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: EISHINGDRELO, Haifeng
APPLICANT: CAI, Jidong
APPLICANT: ARDATI, Mohamad Ali
                                                                                                                                                                                                                TYPE: DNA
ORGANISM: Artificial
                                                                                                                                                                                           OTHER INFORMATION: primer
                                                                                                                                                                                                               FEATURE:
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                                                Local Similarity 100.0%;
nes 13; Conservative
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94 CCTCAGCAGTGTG 106
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100.0%; Pred. No. 2.2e+04;
                                                                                 2.6%;
                                                               Score 13; DB 16;
; Pred. No. 2.2e+04;
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                                                                            Length 24;
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US-09-866-108-12731/c
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PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00664
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SEQ ID NO 7
LENGTH: 24
TYPE: DNA
ORGANISM: Artificial Sequence
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Publication No. US20040096834A1
GENERAL INFORMATION:
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Best Local Similarity
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Patent No. US20020048800A1
                                             PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00662
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00661
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CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
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CURRENT FILING DATE: 2002-11-19
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TITLE OF INVENTION: MODULATION OF HIP-1 PROTEIN INTERACTOR EXPRESSION
FILE REFERENCE: RTS-0431
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
FILE REFERENCE: AEOMICA-7
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00670
                                                                                                                                                                                            APPLICATION NUMBER: PCT/US01/00668
                                                                                                                                                                                                                      APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                   APPLICATION NUMBER: PCT/US01/00669 FILING DATE: 2001-01-30
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HANZEL, David K
RANK, David R.
CHEN, Wensheng
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Pred. No.
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SEQ ID NO 12731
LENGTH: 25
                                                                                                                 SOFTWARE: Aeom
SEQ ID NO 12732
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PRIOR TILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
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PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
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CURRENT FILING DATE: 2001-05-25
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PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 15752
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ORGANISM: Homo sapiens
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ITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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FILING DATE: 2000-09-21
APPLICATION NUMBER: US 60/266,860
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HANZEL, David K
RANK, David R.
CHEN, Wensheng
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-12733
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PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR PILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
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SEQ ID NO 12733
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CURRENT FILING DATE: 2001-05-25
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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APPLICATION NUMBER: US 60/266,860
FILING DATE: 2001-02-05
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00667
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30 CACCTGCTGCTTC 42
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HANZEL, David
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PRIOR FILING DATE: 2001-01-10
PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 2000-09-21
PRIOR FILING DATE: 2000-09-21
PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Acomica Sequence Listing Engine
SEQ ID NO 12734
LENGTH: 25
TYPE: DNA
ORGANISM: Homo sapiens
US-09-866-108-12734
Search completed: February 3, 2005, 00:11:26 Job time : 276.669 secs
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CURRENT APPLICATION NUMBER: US/09/866,108

CURRENT FILING DATE: 2001-05-25

PRIOR APPLICATION NUMBER: US 60/207,456

PRIOR FILING DATE: 2000-05-26

PRIOR APPLICATION NUMBER: GB 24263.6

PRIOR APPLICATION NUMBER: GB 0/236,359

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR APPLICATION NUMBER: PCT/US01/00666

PRIOR FILING DATE: 2000-10-04

PRIOR APPLICATION NUMBER: PCT/US01/00666

PRIOR FILING DATE: 2001-01-30

PRIOR FILING DATE: 2001-01-30
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Best Local Similarity 100.0%; Pred. No. 2.2e+04;
Matches 13; Conservative 0; Mismatches 0;
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APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharrc
APPLICANT: HANZEL, David
APPLICANT: RANK, David
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ITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
ILE REFERENCE: AEOMICA-7
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APPLICATION NUMBER: PCT/US01/00669
FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00663
FILING DATE: 2001-01-30
ADDITION NUMBER: 2001-01-30
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HANZEL, David K.
RANK, David R.
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RESULT 1
US-09-657-453A-51/c
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Patent No. 6458591
GENERAL INFORMATION:
APPLICANT: Brett P. Monia
APPLICANT: Jacqueline Wyatt
TITLE OF INVENTION:
ANTISENSE MODULATION OF PHOSPHORYLASE KINASE ALPHA 2
FILE REFERENCE: RTS-0136
CURRENT APPLICATION NUMBER: US/09/657,453A
CURRENT FILING DATE: 2000-09-07
NUMBER OF SEQ ID NOS: 105
SEQ ID NO 51
LENGTH: 20
         TYPE: DNA
ORGANGE: Artificial Sequence
FEATURE:
OTHER INFORMATION: Antisense Oligonucleotide
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APPLICANT: Lerman, Michael I.

APPLICANT: Minna, John D.

APPLICANT: Minna, John D.

APPLICANT: Hatif, Farida

APPLICANT: Sekido, Yoshitaka

APPLICANT: Sekido, Yoshitaka

APPLICANT: Gao, Boning

APPLICANT: Duh, Fuh-Mei

TITLE OF INVENTION: Calcium Channel Compositions and Methods of Use Thereof

FILE REFERENCE: NIH-05043

CURRENT APPLICATION NUMBER: US/09/470,443

CURRENT FILING DATE: 1999-12-22

EARLIER FILING DATE: 1999-12-30

NUMBER OF SEQ ID NOS: 114

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 37

LENGTH: 22

TYPE: DNA

ORGANISM: Artificial Sequence
APPLICANT: Lake, Staffan

APPLICANT: Wasserman, Wyeth

IIILE OF INVENTION: NOVEL RESPONSE ELEMENT

FILE REFERENCE: 13425-032001

CURRENT FILING DATE: 2000-08-24

PRIOR APPLICATION NUMBER: SE 9904269-9

PRIOR PILING DATE: 1999-11-25

PRIOR PILING DATE: 1999-11-25

PRIOR PILING DATE: 1999-08-31

PRIOR PILING DATE: 1999-08-31

PRIOR APPLICATION NUMBER: SE 9903009-0

PRIOR PILING DATE: 1999-08-26

NUMBER OF SEQ ID NOS: 34

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 12

LENGTH: 25

TYPE: DNA

ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                      US-08-585-684B-790
                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 11; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS:
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/585,60
FILING DATE: January 16, 1996
PRIOR APPLICATION DATA:
APPLICATION UMBER: 60/000,951
APPLICATION UMBER: 50/000,951
FILING DATE: July 7, 1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     tent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 21
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Stinchcomb, Daniel T. APPLICANT: Jarvis, Thale APPLICANT: McSwiggen, James
                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: METHOD AND REAGENT FOR THE TITLE OF INVENTION: INDUCTION OF GRAFT TOLERANCE AND REVERSAL OF IMMUNE RESPONSES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  y Match
3.0%; Score 15; DB 4; 1
Local Similarity 100.0%; Pred. No. 4.6e+02
hes 15; Conservative 0; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                          LENGTH:
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5877021
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                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   UMBER: US/08/585,684B
January 16, 1996
                                                                                                                                                                                                               2.8%;
78.6%;
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                                                                                                                                                                                                               Score 14; DB 2;
Pred. No. 1.5e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/585,684B
FILING DATE: January 16, 1996
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 60/000,951
FILING DATE: July 7, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/078
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Stinchcomb,
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APPLICANT: Stinch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: 67-3510 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44
MEDIUM TYPE: storage
                                                                                                                                                                         APPLICANT: Jarvis, Thale
APPLICANT: McSwiggen, James
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: INDUCTION OF GRAFT TOLERANCE
TITLE OF INVENTION: AND REVERSAL OF IMMUNE RESPONSES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: McSwiggen, TITLE OF INVENTION: ME TITLE OF INVENTION: IN TITLE OF INVENTION: AN
                                                                                                                     NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Ly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 633 West Fifth Street
STREET: 637 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM COMPONERATING SYSTEM:
                                                                                 STREET: 633 West Fifth
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ENGTH:
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                                               Los Angeles
: California
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                                                                                                                                                                                                                                               Stinchcomb, Daniel T. Jarvis, Thale
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IBM Compatible
SYSTEM: IBM P.C. DOS 5.0
                                                                                                                     Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.8%;
78.6%;
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                                                                                                    Street
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Pred. No. 1.5e+03
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Length 15; Indels

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Gaps

COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage

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SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
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APPLICANT: Stinchcomb, Daniel T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                tent No.
                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
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FILING DATE: January 16, 1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/000,951
FILING DATE: July 7, 1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/078
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
                                                                                                                                                             SOFTWARE: FastSEQ Version 1.5 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Jarvis, Thale
APPLICANT: McSwiggen, James
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: INDUCTION OF GRAFT TOLERANCE
TITLE OF INVENTION: AND REVERSAL OF IMMUNE RESPONSES
                                                                                                  PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                    CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. DO SOFTWARE: FASTSEQ Version 1.5
                                                                                 APPLICATION NUMBER:
                                                                                                                        FILING DATE:
                                                                                                                                        APPLICATION NUMBER:
                                                                                                                                                                                               COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C.
                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (213) 489-160
TELEFAX: (213) 955-0440
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Pred. No. 1.5e+03;
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US-09-038-073-791
                                                                                                                                             US-09-038-073-791
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Query Match 2.8%;
Best Local Similarity 78.6%;
Matches 11; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 791,
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APPLICANT: Stinchoomb, Daniel T.
APPLICANT: Jarvis, Thale
APPLICANT: McSwiggen, James
                                                                                                                                                                                                                                                                                                                                                TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: 67-3510 INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
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FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM: IBM P.C. DOS 5.0 SOFTWARE: FastSEQ Version 1.5 CURRENT APPLICATION DATA:
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MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
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Local Similarity 78.6%;
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER:
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ER: 218/078
   Score 14; DB 3; Length 15;
Pred. No. 1.5e+03;
3; Mismatches 0; Indels
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Pred. No. 1.5e+03;
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US-09-549-564-10/c
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; Patent No. 6407284
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                 GENERAL INFORMATION:
APPLICANT: KUDO, JUNKO et al.
TITLE OF INVENTION: METHOD OF RESOLVING 2-OXOBICYCLO[3.1.0]HEXANE-6-CARBOXYLIC
TITLE OF INVENTION: DERIVATIVES
FILE REFERENCE: 2185-0425P
                                                                                                                                                                                                                                                                       Matches
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CURRENT APPLICATION NUMBER: US/09/549,564
                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: FASTSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/038,
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: METHOD AND REAGENT FOR THE TITLE OF INVENTION: INDUCTION OF GRAFT TOLERANCE AND REVERSAL OF IMMUNE RESPONSES
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les 11; Conserv
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STRANDEDNESS: single
TOPOLOGY: linear
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REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 21
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MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
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633 West Fifth Street
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78.6%;
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Pred. No. 1.5e+03;
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CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
                                                                                            ; ORGANISM: Homo sapiens
US-09-866-108A-7839
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SEQ ID NO 7839
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NUMBER OF SEQ ID NOS: 11
SOFTWARE: PatentIn version 3.0
SEQ ID NO 10
                         Query Match
Best Local Similarity
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APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharro
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
                                                                                                                                                                                                                                                                                           PRIOR FILING DATE: 2001-01-30
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                                                                                                                                       TYPE: DNA
                                                                                                                                                            LENGTH:
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APPLICATION NUMBER: PCT/US01/00668
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b. 6686188
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      13;
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CHEN, Wensheng
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    Conservative
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2.6%; Score 13; DB 4;
100.0%; Pred. No. 4.9e+03
tive 0; Mismatches 0
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Pred. No. 1.5e+(
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Sequence 7841, Application US/09866108A
Patent No. 6686188
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SEQ ID NO 7840
                                                                                                                                                             GENERAL
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APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE
FILE REFERENCE: AEOMICA-7
                                                                                                     APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron
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PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/866,108A CURRENT FILING DATE: 2001-05-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Acomica Sequence Listing Engine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
ORGANISM: Homo sapiens
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ITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00664
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                                                                                                                                                             INFORMATION:
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o. 6686188
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                                                  HANZEL, David K
RANK, David R.
CHEN, Wensheng
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HANZEL, David
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                  GENE
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                  EXPRESSED
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                  IN HUMAN
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                  HEART AND MUSCLE
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US-09-866-108A-7841
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN
FILE REFERENCE: ABOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
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                                                           PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/USO1/00666
PRIOR APPLICATION NUMBER: PCT/USO1/00667
PRIOR APPLICATION NUMBER: PCT/USO1/00667
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00664
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00669
PRIOR APPLICATION NUMBER: PCT/USO1/00669
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SEQ ID NO 7841
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Best Local
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CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION UNMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
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PRIOR FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00668
FILING DATE: 2001-01-30
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
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o. 6686188
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Similarity 100.0%;
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RANK, David R.
CHEN, Wensheng
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PENN, Sharron
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Pred. No.
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PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30

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PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR PILING DATE: 2001-05-25
PRIOR PELLING DATE: 2000-05-26
PRIOR PELLING DATE: 2000-05-26
PRIOR PELLING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR PILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
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; ORGANISM: Homo sapiens
US-09-866-108A-7842
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SEQ ID NO 7842
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Patent No. 6686188
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                                                                                                                                                                                                     Remaining Prior Application data removed - NUMBER OF SEQ ID NOS: 15755
SOFTWARE: Acomica Sequence Listing Engine
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CURRENT FILING DATE: 2001-05-25
                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
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APPLICANT: JI, Yonggang
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                                                                                                    ORGANISM: Homo sapiens
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APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
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13; Conserv
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HANZEL, David K.
RANK, David R.
CHEN, Wensheng
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                       100.0%;
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100.0%; Pred. No.
                       2.6%; Score 13; DB 4; Lo
00.0%; Pred. No. 4.9e+03;
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Query Match

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RESULT 17
US-09-422-978-11752/c
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LENGTH: 18
TYPE: DNA
ORGANISM: Homo Sapiens
PEATURE:
NAME/KEY: primer_bind
LOCATION: 1..18
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                                                                                                                                                          NUMBER OF SEQ ID NOS: 11796
SEQ ID NO 11752
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Patent No. 6537751
GENERAL INFORMATION:
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APPLICANT: Blumenfeld, Marta
APPLICANT: Blumenfeld, Marta
APPLICANT: Chumakov, Ilya
APPLICANT: Chumakov, Ilya
TITLE OF INVENTION: Biallelic markers for use in constructing a high density...
FILE REFERENCE: GENSET 020CP1
CURRENT APPLICATION NUMBER: US/09/422,978
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: US 09/298,850
EARLIER APPLICATION NUMBER: US 60/109,732
EARLIER FILING DATE: 1998-11-23
EARLIER APPLICATION NUMBER: US 60/109,732
EARLIER APPLICATION NUMBER: US 60/109,732
EARLIER FILING DATE: 1998-04-21
EARLIER FILING DATE: 1998-04-21
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                                                                                                                                                                                                                                           FILE REFERENCE: GENSET.020CP1
CURRENT APPLICATION NUMBER: US/09/422,978
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: US 09/298,850
EARLIER FILING DATE: 1999-04-21
EARLIER FILING DATE: 1998-11-23
EARLIER FILING DATE: 1998-11-23
                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Cohen, Daniel
APPLICANT: Blumenfeld, Marta
APPLICANT: Chumakov, Ilya
TITLE OF INVENTION: Biallelic markers for use
                                                                                                                                                                                                  EARLIER APPLICATION NUMBER: US 60/082,614 EARLIER FILING DATE: 1998-04-21
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NAME/KEY: primer_bind
LOCATION: 1..18
OTHER INFORMATION: downstream
                                                                                      TYPE: DNA
ORGANISM: Homo Sapiens
                                                                                                                                       ENGTH: 18
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      amplification
    primer 99-5075
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PatentIn Release #1.0, Version #1.30

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US-09-544-398B-478
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; ORGANISM: Homo sapiens
US-09-544-398B-478
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Best Local :
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SEQ ID NO 478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent No. 67704
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
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PRIOR APPLICATION NUMBER: US 09/229,319
PRIOR FILING DATE: 1999-01-13
PRIOR APPLICATION NUMBER: US 60/071,449
PRIOR FILING DATE: 1998-01-13
PRIOR APPLICATION NUMBER: US 60/105,511
PRIOR FILING DATE: 1998-10-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Carulli, John P.
APPLICANT: Little, Randall D.
APPLICANT: Recker, Robert R.
                                                                                                                                                                                                                                                                                                                                                                                 atent No. 5693473
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/544,398B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Johnson, Mark L.
TITLE OF INVENTION: High bone mass gene of 11q13.3
FILE REFERENCE: 032796-013
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                        APPLICANT: APPLICANT:
                             COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                          TITLE OF INVENTION:
                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                         APPLICANT:
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OPERATING SYSTEM:
                                                              COUNTRY: U
                                                                                                                                            ADDRESSEE:
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                                                                                                         Washington
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                                                                                                                                                                                                                                                                                                                                                                                                                Application US/08480784
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ACAAGTGTCTGGA 14
                                                                                                                         E: Venable, Baetjer, Howard & Civiletti, LLP
1201 New York Avenue, N.W., Suite 1000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Application US/09544398B
                                                                               USA
                                                                                                                                                                                   Futreal, P. Andrew
VENTION: 17q-Linked Breast and Ovarian Cancer
VENTION: Susceptibility Gene
                                                                                                                                                                                                                                   Tavtigian, Sean V
Wiseman, Roger W.
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Goldgar, David E.
Miki, Yoshio
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                I: Floppy disk
IBM PC compati
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 PC-DOS/MS-DOS
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100.0%;
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; Pred. No. 4.9e+
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RESULT 20
US-08-483-553-4/c
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Best Local :
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                                                                                                                                                                                     APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                     APPLICANT:
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LENGTH: 20 base pairs
TYPE: nucleic acid
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ORIGINAL SOURCE:
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REFERENCE/DOCKET NUMBER: 24
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-962-4810
TELEFAX: 202-962-8300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 16-SEP-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/300,266
FILING DATE: 02-SEP-1994
PRIOR APPLICATION DATA:
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PRIOR APPLICATION NUMBER: US 08/409,305
APPLICATION NUMBER: US 08/409,305
                                                CORRESPONDENCE ADDRESS: ADDRESSEE: Venable,
                                                                                                 APPLICANT: Wiseman, Roger W.
APPLICANT: Futreal, P. Andrew
TITLE OF INVENTION: 17q-Linked Breast and Ovarian Cancer
TITLE OF INVENTION: Susceptibility Gene
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FILING DATE: 29-NOV-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/308,104
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PRIOR APPLICATION DATA:
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                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                      APPLICANT:
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                                 STREET:
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Washington
: DC
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                                 E: Venable, Baetjer, Howard & Civiletti, 1201 New York Avenue, N.W., Suite 1000
                                                                                                                                                                    Tavtigian, Sean V.
                                                                                                                                                                                   Harshman, Keith D. Shattuck-Eidens, D
                                                                                                                                                                                                                                                        Miki, Yoshio
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Goldgar, David E.
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12-AUG-1994
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US-08-487-002-4/c
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Patent No. 5710001
                                                                                                                                                                                                                                                                                                                                                                                             Query Match 2.6%; So
Best Local Similarity 100.0%; P
Matches 13; Conservative 0;
                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Shattuck-Eidens, Donna
APPLICANT: Simard, Jacques
APPLICANT: Emi, Mitsuru
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 202-962-8300 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORIGINAL SOURCE:
ORGANISM: Homo sapiens
IMMEDIATE SOURCE:
        APPLICANT: Nakamura, Yusuke
APPLICANT: Durocher, Francine
TITLE OF INVENTION: 17q-Linked Breast and Ovarian Cancer
TITLE OF INVENTION: Susceptibility Gene
NUMBER OF SEQUENCES: 85
CORRESPONDENCE ADDRESS:
ADDRESSEE: Venable, Baetjer, Howard & Civiletti, LLP
STREET: 1201 New York Avenue, N.W., Suite 1000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 12-AUG-1994
ATTORNEY/AGENT INFORMATION:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/300,266
FILING DATE: 02-SEP-1994
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08
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APPLICATION NUMBER: US 08/409,305
FILING DATE: 24-MAR-1995
PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER: 28,957
REFERENCE/DOCKET NUMBER: 24
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OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
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; Pred. No.
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US-08-483-554B-4/c
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                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Skolnic
APPLICANT: Goldgar
APPLICANT: Miki, )
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INFORMATION FOR SEQ ID NO:
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APPLICATION NUMBER: US 08/
FILING DATE: 02-SEP-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
FILING DATE: 12-AUG-1994
                                                                                 APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Homo sapiens
IMMEDIATE SOURCE:
CLONE: 8754 B
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/487,002
FILING DATE:
CLASSIFICATION: 424
                                                   APPLICANT:
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REGISTRATION NUMBER: 28,957
REFERENCE/DOCKET NUMBER: 241
TELECOMMUNICATION INFORMATION:
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PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/308,104
FILING DATE: 16-SEP-1994
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PRIOR APPLICATION DATA:
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TO TO NO:
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                                                                                                                                               Skolnick, Mark
Goldgar, David
                                            Harshman, Keith D.
Shattuck-Eidens, Donna M.
Tavtigian, Sean V.
                                                                                                               Swenson, Jeff
                               Wiseman, Roger W.
                                                                                                  Kamb, Alexander
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17q-Linked Breast and Ovarian Cancer
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US-08-488-011B-4/c
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                                                          Sequence 4, Application US/08488011B
Patent No. 5753441
GENERAL INFORMATION:
APPLICANT: Skolnick, Mark H.
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Best Local S
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               APPLICANT:
APPLICANT:
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MEDIUM TYPE: Ploppy disk
COMPUTER: PLOPPY DOSPATIBLE
COMPUTER: PM PC COMPUTED:
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
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HYPOTHETICAL: N
ORIGINAL SOURCE:
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ADDRESSEE: Venable, Baetjer, Howard & Civiletti, LLP STREET: 1201 New York Avenue, N.W., Suite 1000
                                                                                                                                                                                                                                                                                                                                                      IMMEDIATE SOURCE:
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LENGTH: 20 base pairs
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PRIOR APPLICATION NUMBER: US 08/308,104
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PRIOR APPLICATION DATA:
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APPLICATION NUMBER: US 08/409,305
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Ihnen, Jeffrey L. REGISTRATION NUMBER: 28,957
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CLASSIFICATION:
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                            Skolnick, Mark I
Goldgar, David I
Miki, Yoshio
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Kamb, Alexander
             Swenson, Jeff
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US-08-488-011B-4
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Best Local
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APPLICATION NUMBER: US
FILING DATE: 24-MAR-1s
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US
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HYPOTHETICAL: NO
ORIGINAL SOURCE:
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FILING DATE: 16-SEP-1994
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 08/300,266
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/
FILING DATE: 07-JUN-199
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APPLICATION NUMBER: 1
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PRIOR APPLICATION UNMBER: US 08/308,104
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TELECOMMUNICATION INFORMATION:
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CTTY: Washington
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                                                                                                                                                            ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                    NAME: Ihnen, Jeffrey L. REGISTRATION NUMBER: 28
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ZIP: 20005
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                          432 CATTCCTGCTTCC 444
16 CATTCCTGCTTCC
                                                        13;
                                                                      Similarity
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Shattuck-Eidens, I
                                                        Conservative
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SYSTEM: PC-DOS/MS-DOS
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Susceptibility Gene
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Pred. No.
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RESULT 24 US-08-850-727-4/c

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Sequence 4, Application US/08850727 Patent No. 6162897

ENERAL INFORMATION:

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Matches 13
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HYPOTHETICAL: NO
ORIGINAL SOURCE:
ORGANISM: Homo (
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                                                                                                                                                           IMMEDIATE SOURCE:
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LENGTH: 20 base pairs
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APPLICATION NUMBER: US 08
FILING DATE: 02-SEP-1994
PRIOR APPLICATION DATA:
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PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/308,104
FILING DATE: 16-SEP-1994
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APPLICATION NUMBER: US 08/483,554
FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Futreal, P. Andrew
TITLE OF INVENTION: 17q-Linked Breast and Ovarian Cancer
TITLE OF INVENTION: Susceptibility Gene
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                                                                                                                                                                                                                                                TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                               CLONE: 8754 B
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                               432 CATTCCTGCTTCC 444
16 CATTCCTGCTTCC 4
                                                              l Similarity
13; Conserv
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1201 New York Avenue,
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                                                            2.6%; Score 13; DB.larity 100.0%; Pred. No. 4.:
Conservative 0; Mismatches
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Wiseman, Roger W.
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rk Avenue, N.W., Suite 1000
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                                                                             DB 3; Lo
                                                                                           Length 20
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; OTHER INFORMATION: Description of Artificial Sequence: primer reverse ; OTHER INFORMATION: for D17S1322 US-09-445-174B-2
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                                                                                                                                               SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 2
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FILE REFERENCE: 03715.0048-00000
CURRENT APPLICATION NUMBER: US/09/230,652A
CURRENT FILING DATE: 1999-05-17
EARLIER APPLICATION NUMBER: FR 96 09733
EARLIER APPLICATION NUMBER: FR 97 04680
EARLIER FILING DATE: 1997-04-16
EARLIER APPLICATION NUMBER: FR 97 04680
EARLIER FILING DATE: 1997-04-16
EARLIER APPLICATION NUMBER: PCT/FR97/01433
EARLIER FILING DATE: 1997-07-31
NUMBER OF SEQ ID NOS: 133
SOPTMARE: Patentin Ver: 2.1
                                                                                                                                                                                         PRIOR APPLICATION NUMBER: EP 97201700.8 PRIOR FILING DATE: 1997-06-04 NUMBER OF SEQ ID NOS: 23
                                                                                                                                                                                                                                                                                                                                         APPLICANT: Devilee, Peter
TITLE OF INVENTION: A diagnostic test kit for determining
TITLE OF INVENTION: for breast and ovarian cancer, materi
TITLE OF INVENTION: for such determination
FILE REFERENCE: 294-78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Joutel, Anne
APPLICANT: Bousser, Marie-Germaine
APPLICANT: Bach, Jean-Francois
                                                                                                                                                                                                                                                        CURRENT FILING DATE: 2001-06-11
PRIOR APPLICATION NUMBER: PCT/NL98/00325
PRIOR FILING DATE: 1998-06-03
                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/445,174B
CURRENT FILING DATE: 2001-06-11
                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                      LENGTH: 20
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: van Ommen, Garrit J.B.
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                                                            ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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4.9e+03;
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Query Match Best Local Similarity

Score 13; Pred. No.

DB 4; Length 20; . 4.9e+03;

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RESULT 27
PCT-US95-10202-4/c
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                                                                                                                                                TELEFAX: 202-962-8300
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Nakamura, Yusuke
APPLICANT: Durocher, Francine
TITLE OF INVENTION: In the 170
TITLE OF INVENTION: Susceptible
NEWTON OF INVENTION: Susceptible
IMMEDIATE SOURCE:
CLONE: 8754 B
                                                ORIGINAL SOURCE
                                                                MOLECULE TYPE: DI
HYPOTHETICAL: NO
                                                                                                                                                                                                                  NAME: Ihnen, Jeffrey L.
REGISTRATION NUMBER: 28,957
REFERENCE/DOCKET NUMBER: 24
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-962-4810
                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 0
FILING DATE: 12-AUG-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/348,824
PTITUG DATE: 29-NOV-1994
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PRIOR APPLICATION DATA:
US
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 02-SEP-
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 07-JUN-
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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APPLICATION NUMBER: 1
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 1201 No.
                               ORGANISM: Homo sapiens
                                                                                                             STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 08-308,104 FILING DATE: 16-SEP-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
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FILING DATE: 07-JUN-1995
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1201 New York Avenue, N.W., Suite 1000
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                                                                                                    linear
                                                                                                                                                                                                                                                      INFURE 1. Jeffrey L. 28,957
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                                                                                 DNA (genomic)
                                                                                                               single
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                                                                                                                                                                                                                                                                                                                                     US 08/289,221
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Best Local :
                                                TELEFAX: 202-96 INFORMATION FOR SEQ
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                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/300,266
FILING DATE: 02-SEP-1994
                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 07-UUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/409,305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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PRIOR APPLICATION NUMBER: US 08/348,824
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                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
APPLICATION NUMBER: PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                               FILING DATE: 12-AUG-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                PRIOR APPLICATION DATA:
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                                                                                               TELECOMMUNICATION INFORMATION:
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                                                                                                                              NAME: Ihnen, Jeffrey L. REGISTRATION NUMBER: 28
                                                                              TELEPHONE:
                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
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STREET: 12
                                                                                                                 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.30
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                                                                  202-962-8300
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Miki, Yoshio
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                                                                                 202-962-4810
                                                                                                                                                                                                                                                                                  16-SEP-1994
                                                                                                                                                                                                                                                                                                                                     29-NOV-1994
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100.0%; Pred. No. 4.9e+03
                                                                                                                                                                                                  US 08/289,221
                                                                                                                                                                                                                                                                                                  US 08-308,104
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                                                                                                                             28,957
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                                                                                                                 24884-109347
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                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08-308,104
FILING DATE: 16-SEP-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/300,266
FILING DATE: 02-SEP-1994
PRIOR APPLICATION DATA:
APPLICATION UMBER: US 08/289,221
APPLICATION UMBER: US 08/289,221
FILING DATE: 12-AUG-1994
ATTORNEY/AGENT INFORMATION:
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APPLICANT: Futreal,
TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OKCANISM: Homo sapiens
IMMEDIATE SOURCE:
                                                                                                                                                                                                                APPLICATION NUMBER: US 08/409,305
FILING DATE: 24-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/348,824
FILING DATE: 29-NOV-1994
                                                                                                                                                                                                                                                                                   PILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA:
ADDITICATION NUMBER: US 08/409,305
                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
CRASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: FC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
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HYPOTHETICAL: NO
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TOPOLOGY: lin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       432 CATTCCTGCTTCC 444
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16 CATTCCTGCTTCC 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  E: Venable, Baetjer, Howard & Civiletti, LLP 1201 New York Avenue, N.W., Suite 1000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Futreal, P. Andrew
VENTION: Method for Diagnosing a
VENTION: Predisposition for Breast and Ovarian Cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kamb, Areasseth D. Harshman, Keith D. Harshman, Keith D. Harshman, M. Harshman, Donna M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Skolnick, Mark H. Goldgar, David E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Wiseman, Roger W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Shattuck-Eidens, D
Tavtigian, Sean V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Swenson, Jeff
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA (genomic)
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100.0%; Pred. No. 4.
tive 0; Mismatches
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CRIGINAL SOURCE:
CRGANISM: Homo:
IMMEDIATE SOURCE:
CLONE: 8754 B
PCT-US95-10220-4
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                                                                                                                     Sequence 5, Application US/09587945 Patent No. 6709860 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
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                  APPLICANT: Carlsson, Peter
TITLE OF INVENTION: Animal Model
FILE REFERENCE: 10806-117A
CURRENT APPLICATION NUMBER: US/09/587,945
CURRENT FILING DATE: 2000-06-06
                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: GB
PRIOR FILING DATE: 1998-11-11
PRIOR APPLICATION NUMBER: PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR FILING DATE:
 PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 24884-109347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
                                                                                                                                                                                                                                                                          127 TGCCACCTGTACT 139
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                                                                                                      Enerback,
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US 60/190,692
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CURRENT FILING DATE: 2000-07-14

PRIOR APPLICATION NUMBER: GB 9801391

PRIOR APPLICATION NUMBER: GB 9801391

PRIOR APPLICATION NUMBER: GB 9801391

PRIOR APPLICATION NUMBER: GB 9801391
                                                             ; SEQ ID NO 2
; LENGTH: 22
; TYPE: DNA
; ORGANISM: Human interleukin-4 promoter
US-09-617-548-2
                                                                                                                                                                                                                                                                                                                                                                         Sequence 2, Application US/09617548 Patent No. 6476214 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION: TELEPHONE: 202-962-4810
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                                                                                                                                                                             1999-01-20
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100.0%; Pred. No. 4.9e+03;
tive 0; Mismatches 0;
             2.6%; Score 13;
100.0%; Pred. No.
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0,
Mismatches
                4.9e+03
                            Length 22
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PRIOR FILING DATE:

2000-03-20

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RESULT 32
US-08-808-550-8/c
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                                                                       US-08-808-550-8
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Patent No. 5871992
GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 5
LENGTH: 22
 Matches
                                  Query Match
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PRIOR FILING DATE: 1997-05-26
NUMBER OF SEQ ID NOS: 26
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                          TELEFAX: 201-343-1684
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
ADDRESSEE: David A. Jackson, Esq.
STREET: 411 Hackensack Ave, Continental Plaza,
                                                                                                      MOLECULE TYPE: other nucleic acid DESCRIPTION: /desc = "Primer P3"
                                                                                                                                                                                                                                                               TELEPHONE: 201-487-5800
                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/808,550
                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
ODERATING SYSTEM: PC-DOS/MS-DOS
ODERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: FLOCE
CITY: Hackensack
CTATE: New Jersey
                                                                                       HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Teebor, George W.
APPLICANT: Hilbert, Timothy P.
TITLE OF INVENTION: MAMPALIAN ENDONUCLEASE III AND
TITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC USES THEREOF
                  Local Similarity
                                                                                                                                                                                                                                                                                                NAME: Jackson Esq., David A.
REGISTRATION NUMBER: 26,742
REFERENCE/DOCKET NUMBER: 1049-1-001 N
                                                                                                                                         TOPOLOGY:
                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/0
FILING DATE: 26-FEB-1997
                                                                                                                                                                                            ENGTH:
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                  2.6%; Score 13; DB 2;
100.0%; Pred. No. 4.9e+C
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; Pred. No. 4.9e+03;
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 Mismatches
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                4.9e+03;
                                  Length 24;
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Indels
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US-08-859-998-767
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US-08-691-814B-53/c
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                                                                                                              Patent No. 599407
                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 13; Conserv
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GENERAL INFORMATION:
                                                                                                                              Sequence 767, Application US/08859998
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                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 202-371-2543
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
APPLICANT: Chenchik, Alex
APPLICANT: Jokhadze, George
APPLICANT: Bibilashvilli, Robert
TITLE OF INVENTION: METHED OF ASSAYING DIFFERENTIAL
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 1375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Basset, Paul
APPLICANT: Byrne, Jennifer
TITLE OF INVENTION: Isolated Nucleic Acid Molecules Useful
TITLE OF INVENTION: as Leukemia Markers and in Breast Cancer Prognosis
NUMBER OF SEQUENCES: 124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Rio, Marie-Christine
APPLICANT: Tomasetto, Catherine
APPLICANT: Basset, Paul
APPLICANT: Byrne, Jennifer
                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 1100 New CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
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                                                                                                                                                                                                                                             138 CTGGGGCTGCACC 150
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20005-3934
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                                                                                                                                                                                                                                                                                Conservative
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100.0%; Pred. No.
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JS-09-225-___
JS-09-225-___
Sequence 767, Ap.
Patent No. 6352829
Patent INFORMATION:
GENERAL INFORMATION:
APPLICANT: Chenchik, Alex
Jokhadze, George
Bibilabilii, Rob
Bibilabilii, Rob
O'N'TION: METHOD O'
TYPRESS'
                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 35
US-09-225-928-767
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 24 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 09
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-322-5070
TELEPAX: 415-854-0875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY AGENT INFORMATION:
NAME: Field Bret E.
REGISTRATION NUMBER: 37,620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Fish & Richardson, P.C. STREET: 2200 Sand Hill Road, Suite 100 CITY: Menlo Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: oligonucleotide primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/01
FILING DATE: 21-MAY-1997
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ZIP: 94025
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                  COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/09/225,928
FILING DATE: 05-Jan-199
CLASSIFICATION: Unknown>
                                                                                                                                       ZIP: 94025
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSE: Fish & Richardson, P.C.
STREET: 2200 Sand Hill Road, Suite 100
CITY: Menlo Park
STATE: CA
       PRIOR
                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: METHOD OF ASSAYING DIFFERENTIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16 AGCCACGCCCTGT 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13;
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                                                                                                                                                                                               COUNTRY: US
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APPLICATION DATA
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100.0%; Pred. No. 4.1
tive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                   EXPRESSION
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                                                                                          2.0
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US-09-225-201B-767
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                TYPE: nucleic acid
STRANDEDNESS: single
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OTHER INFORMATION: oligonucleotide primer; SEQUENCE DESCRIPTION: SEQ ID NO: 767: US-09-225-928-767
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GENERAL INFORMATION:
                                              TELEFAX: 415-854-0875
INFORMATION FOR SEQ ID NO: 767:
SEQUENCE CHARACTERISTICS:
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INFORMATION FOR SEQ ID NO: 767:
                                                                                                                                                                      APPLICATION NUMBER: US/08/859,998
FILING DATE: 21-MAY-1997

ATTORNEY AGENT INFORMATION:
NAME: Field, Bret E.
REGISTRATION NUMBER: 37,620
REGISTRATION NUMBER: 09096/00200
REFERENCE/DOCKET NUMBER: 09096/00200
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-322-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette

COMPUTER: IBM Compatible

OPERATING SYSTEM: Windows 95

SOFTWARE: FastSEQ for Windows Version 2.0

CURRENT APPLICATION DATA:

APPLICATION UNMBER: US/09/225,201B

FILING DATE: 05-Jan-199

CLASSIFICATION: VUMCHOWN>

PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 08/859,998
FILING DATE: 21-MAY-1997
ATTORNEY_AGENT INFORMATION:
NAME: Field, Bret E.
REGISTRATION NUMBER: 37,620
REGISTRATION NUMBER: 09096/01
TELEPHONE: 415-322-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 2200 Sand Hill Road, Suite 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bibilashvilli, Robe
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Menlo Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
LENGTH: 24 base pairs
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Sequence 17, Application US/09667135

PARTENT NO. 6521749

GENERAL INFORMATION:

APPLICANT: Vincent Ling
APPLICANT: Kyriaki Dunussi-Joannopoulos

TITLE OF INVENTION: NOVEL GL50 MOLECULES AND USES THEREFOR
FILE REFERENCE: GNN-007

CURRENT APPLICATION NUMBER: US/09/667,135

CURRENT APPLICATION NUMBER: US/09/667,135

CURRENT FILING DATE: 2000-09-21

NUMBER OF SEQ ID NOS: 38

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 17

LENGTH: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Sonigo, Pierre
APPLICANT: Sonigo, Pierre
APPLICANT: Sonigo, Pierre
APPLICANT: Sonigo, Christian
APPLICANT: Brechot, Christian
APPLICANT: Brechot, Christian
APPLICANT: Courgnard, Valerie
TITLE OF INVENTION: OLIGONUCLEOTIDE SEQUENCES FOR THE AMPLIFICATION OF THE
TITLE OF INVENTION: AND THEIR USES FOR IN VITRO DIAGNOSIS OF THE INFECTIONS
TITLE OF INVENTION: DUE TO THESE VIRUSES
FILE REFERENCE: 2356.0065-01
CURRENT APPLICATION NUMBER: US/08/343,998A
CURRENT APPLICATION NUMBER: 07/820,600
EARLIER APPLICATION NUMBER: 07/820,600
EARLIER APPLICATION ONTHE : 1992-01-22
EARLIER APPLICATION ONTHE : 1990-06-05
NUMBER OF SEO ID NOS: 25
CORTAINER APPLICATION ONTHE : 1990-06-05
NUMBER OF SEO ID NOS: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; SEQUENCE DESCRIPTION: SEQ ID NO: 767: US-09-225-201B-767
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; OTHER INFORMATION: primer US-09-667-135-17
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US-09-667-135-17
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SEQ ID NO 12
LENGTH: 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 2.6
Best Local Similarity 100
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                               TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Human immunodeficiency virus type
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         124 TTCTGCCACCTGT 136
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100.0%; Pred. No. 4.9e+03;
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100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                               ; ORGANISM: Homo sapiens US-09-866-108A-12731
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; Sequence 12731, Application US/09866108A
                                                                                                                                                 US-09-866-108A-12732/c
                                                                                                                                                                                                                                                                                                                                                                                                                                        ; Patent No. 66861
; SEQ ID NO 12731
; LENGTH: 25
; TYPE: DNA
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Best Local S
Matches 13
Sequence 12732, Application US/09866108A
Patent No. 6686188
GENERAL INFORMATION:
APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
                                                                                                                                                                                                                                                                                                                                      Best
                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
FILE REPERBOCE ABOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR PILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: GB 4263.6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BENERAL INFORMATION:
APPLICANT: GU, Yizhong
APPLICANT: JI, Yongga
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 15755
SOFTWARE: Acomica Sequence Listing Engine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
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PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
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PRIOR APPLICATION NUMBER: PCT/US01/00663
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100.0%; Pred. No.
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L00.0%; Pred. No.
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APPLICANT: CHEN, MERICANT, APPLICANT: SHANNON, MARK

TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN

FILE REFERENCE: AEOMICA-7

CURRENT APPLICATION NUMBER: US/09/866,108A

CURRENT FILLING DATE: 2001-05-25

PRIOR APPLICATION NUMBER: US 60/207,456

PRIOR FILLING DATE: 2000-05-26

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR FILLING DATE: 2000-09-27

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR APPLICATION NUMBER: PCT/US01/00666

PRIOR APPLICATION NUMBER: PCT/US01/00667

PRIOR APPLICATION NUMBER: PCT/US01/00667

PRIOR APPLICATION NUMBER: PCT/US01/00667

PRIOR APPLICATION NUMBER: PCT/US01/00667

PRIOR FILLING DATE: 2001-01-30

PRIOR FILLING DATE: 2001-01-30
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TITLE OF INVENTION: MYCSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION UNUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR APPLICATION NUMBER: PCT/US01/00669
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US-09-866-108A-12733/c
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APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharry
APPLICANT: HANZEL, David
APPLICANT: RANK, David
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Best Local
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Patent No. 6686188
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SOFTWARE: Recmica Sequence Listing Engine
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PRIOR APPLICATION NUMBER: PCT/US01/00663
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ORGANISM: Homo sapiens
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FILING DATE: 2001-01-30
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PENN, Sharron G.
HANZEL, David K.
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100.0%; Pred. No.
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SOPTWARE: Aeomica Sequence
Patent No. 6686188
SEQ ID NO 12734
LENGTH: 25
TYPE: DNA
ORGANISM: Homo sapiens
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; ORGANISM: Homo sapiens
US-09-866-108A-12733
US-09-866-108A-12734
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PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR PELICATION NUMBER: US 60/207,456
PRIOR PELICATION NUMBER: GB 24263.6
PRIOR PELICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR PLING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00667.
PRIOR APPLICATION NUMBER: PCT/US01/00667.
PRIOR APPLICATION NUMBER: PCT/US01/00667.
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR APPLICATION NUMBER: PCT/US01/00664
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NUMBER OF SEQ ID NOS: 15755
SOFTWARE: Acomica Sequence Listing Engine
Patent No. 6686188
SEQ ID NO 1733
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                                                                                                                                                         Remaining Prior Application NUMBER OF SEQ ID NOS: 15755
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APPLICANT: RANK, David R.
APPLICANT: CHANNOW, Warsheng
APPLICANT: SHANNOW, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: ABOMICA-7
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PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR: FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
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PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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APPLICANT: JI, Yonggang
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APPLICATION NUMBER: PCT/US01/00669
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
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Pred. No.
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US-09-866-108A-12736/c
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               GENERAL INFORMATION:
APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggan
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SEQ ID NO 12735
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 APPLICANT:
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PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
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APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                            ENGTH: 25
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00667
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APPLICATION NUMBER: PCT/US01/00663
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FILING DATE: 2001-01-30
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RANK, David R.
CHEN, Wensheng
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Yonggang
N, Sharron
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Pred. No.
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RESULT 45
US-09-866-108A-12737/c
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US-09-866-108A-12736
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR PELICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
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Matches
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Patent No. 6686188
SEQ ID NO 12736
LENGTH: 25
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APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 201-05-25
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
FILE REFERENCE: AECOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
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APPLICANT:
APPLICANT:
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APPLICANT: JI, Yongga
APPLICANT: PENN, Shar
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PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
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PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
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PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
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PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
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PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
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mes 13; Conserv
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HANZEL, David
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; Pred. No.
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4.9e+03
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RESULT 46
US-09-866-108A-12738/c
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; ORGANISM: Homo sapiens
US-09-866-108A-12737
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CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
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Best Local Similarity
                                                           Patent No. 66863
SEQ ID NO 12738
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SEQ ID NO 12737
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APPLICANT: JI, Yongga
APPLICANT: PENN, Shar
                                                                                                         SOFTWARE: Acomica Sequence Listing Engine
                                                                                                                                PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 15755
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
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APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
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ORGANISM: Homo sapiens
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APPLICATION NUMBER: PCT/US01/00665
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CATION NUMBER: PCT/US01/00668
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PENN, Sharron G.
HANZEL, David K.
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ON NUMBER: PCT/US01/00669
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US-09-866-108A-12739
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CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR APPLICATION NUMBER: PCT/US01/00664
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                                                                                                                                                                                                              Patent No. 6686188
SEQ ID NO 12739
LENGTH: 25
TYPE: DNA
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00668
FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00669
FILING DATE: 2001-01-30
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17 CACCTGCTGCTTC 5
                            30 CACCTGCTGCTTC 42
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                                                                                  13; Conservative 0;
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David K.
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100.0%; P7
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                                                                                                                          DB 4; Length 25;
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US-09-866-108A-12740/c
US-09-866-108A-12740, Application US/09866108A
; Patent NO. 6686188
; GENERAL INFORMATION:
; APPLICANT: GU, Yizhong
; APPLICANT: JI, Yonggang

RESULT 48

PENN, Sharron

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US-09-866-108A-12741/c
                               CURRENT APPLICATION NUMBER: US/09/866,108A CURRENT FILING DATE: 2001-05-25 PRIOR APPLICATION NUMBER: US 60/207,456 PRIOR FILING DATE: 2000-05-26 PRIOR APPLICATION NUMBER: GB 24263.6 PRIOR APPLICATION NUMBER: US 60/236,359 PRIOR APPLICATION NUMBER: US 60/236,359 PRIOR FILING DATE: 2000-09-27
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108A-12740
                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 12741, Application US/09866108A Patent No. 6686188
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SEQ ID NO 12740
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PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2000-09-27
                                                                                                                                                                                                             APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
FILE REFERENCE: AEOMICA-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Aeomica Sequence Listing Engine
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APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
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ITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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APPLICATION NUMBER: PCT/US01/00668
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               APPLICATION NUMBER: PCT/US01/00666
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13; Conservative (
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; Sequence 12742, Application US/09866108A
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SEQ ID NO 12742
LENGTH: 25
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SEQ ID NO 12741
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Best Local Similarity
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                                                     Remaining Prior Application data removed - NUMBER OF SEQ ID NOS: 15755
SOFTWARE: Acomica Sequence Listing Engine
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TITLE OF INVENTION: WOSSN-LIKE GENE EXPRES
FILE REFERICE: ABOMICA-T
CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
                                                                                                             PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
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PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
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FILING DATE: 2001-01-30
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APPLICATION NUMBER: US 60/236,359
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                                                                                                                                                                                                                                                  APPLICATION NUMBER: PCT/US01/00669
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INVENTION: MYOSIN-LIKE GENE EXPRESSED
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PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2001-05-25
PRIOR PRIOR PELLOATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR PRIOR PELLOATION NUMBER: PCT/US01/00666
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00664
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                                                                                  RESULT 52
US-08-330-790A-4/c
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; ORGANISM: Homo sapiens
US-09-866-108A-12743
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SEQ ID NO 12743
LENGTH: 25
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Best Local
                                 Sequence 4, Application US/08330790A Patent No. 5629178
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Best Local
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APPLICANT: JI, Yongga
APPLICANT: PENN, Shar
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
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APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
  GENERAL INFORMATION:
APPLICANT: Demers
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13; Conserv
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Demers, Daniel B
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100.0%; Pred. No.
tive 0; Mismatc
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                                                                                                                                                                                                                                                                                                                                                           Sequence 4, Application US/08468658
Patent No. 5656461
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                                                                                                                                                                                                                         APPLICANT: Demers, Daniel
TITLE OF INVENTION: A Met
TITLE OF INVENTION: in th
TITLE OF INVENTION: Emplo
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: MCDERMOTT, M
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TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBÉR OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 435
PRIOR APPLICATION NUMBER:
                           SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/468,65
                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 28,553
REFERENCE/DOCKET NUMBER: 75
TELECOMMUNICATION INFORMATION:
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 28 base par
TYPE: nucleic acid
                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                       STREET: 1850 K St
CITY: Washington
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
FILING DATE: 0 CLASSIFICATION:
                                                                                                                                                             COUNTRY:
                                                                                                                                                                              STATE: D.C.
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             06-JUNE-1995
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in the Polymerase Chain Reaction
Employing Peptide Nucleic Acids (PNA)
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Employing Peptide Nucleic Acids
                              US/08/468,658
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GENERAL INFORMATION:
APPLICANT: Genetics & IVF Institut
TITLE OF INVENTION: A Method for I
TITLE OF INVENTION: in the Polymer
TITLE OF INVENTION: Employing Pept
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Best Local Similarity
Matches 13; Conserv
                                                    Query Match
Best Local :
                                     Matches
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                   TELEPHONE: 202-429-06 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 28 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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APPLICATION NUMBER: US 08/33
FILING DATE: 28-OCTOBER-1994
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                  TELEPHONE: 202-429-0625
                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION: NAME: Paul Devinsky
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Paul Devinsky
REGISTRATION NUMBER: 28
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: FILING DATE:
                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 75
                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
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                                   l Similarity
13; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Genetics & IVF Institute, Inc.
VERVION: A Method for Enhancing Amplification
VERVION: in the Polymersase Chain Reaction
VERVION: Employing Peptide Nucleic Acids (PNA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.6%; Score 13; DB 1; ilarity 100.0%; Pred. No. 4.9e+(Conservative 0; Mismatches
                                   2.6%; Score 13; DB 5; Lilarity 100.0%; Pred. No. 4.9e+03; Conservative 0; Mismatches 0;
                                                                                                                             linear
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4.9e+03;
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                                     Indels
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CURRENT APPLICATION NUMBER: US/08/297,395A
CURRENT FILING DATE: 1994-08-11
EARLIER APPLICATION NUMBER: 08/059,189
EARLIER FILING DATE: 1993-05-06
EARLIER APPLICATION NUMBER: 07/502,559
EARLIER FILING DATE: 1990-03-30
EARLIER APPLICATION NUMBER: PCT/US88/02139
EARLIER APPLICATION NUMBER: PCT/US88/02139
EARLIER FILING DATE: 1988-06-24
EARLIER FILING DATE: 1988-06-24
EARLIER FILING DATE: 1987-06-24
EARLIER FILING DATE: 1987-06-24
EARLIER FILING DATE: 1987-06-24
EARLIER FILING DATE: 1987-06-24
NUMBER OF SEQ ID NOS: 84
SOFTWARE: FastSEQ for Windows Version 3.0
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Best Local Similarity
Matches 13; Conserv
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           COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: #1.30 (EPO)

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/765,340

FILING DATE: 23-DEC-1996

PRIOR APPLICATION DATA:

APPLICATION NUMBER: UP 145146/94

FILING DATE: 27-JUN-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 104,
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APPLICANT: David A. Hafler
TITLE OF INVENTION: EPFILDES DERIVED FROM IMMUNODOMINANT
TITLE OF INVENTION: EPITOPES OF MYELIN BASIC PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 30
TYPE: DNA
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APPLICANT:
APPLICANT:
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ADDRESSEE: MORGAN & FINNEGAN, L.L.P.
                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: AN ANTIS
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PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                           ZIP: 10154
                                                                                                                                                                                                                                                                                                                        CITY: NEW YORK
STATE: NEW YORK
                                                                                                                                                                                                                                                                                                 COUNTRY:
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TANAKA, Y.,
MATSUDA, Y.,
KONDO, S.
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100.0%; Pred. No. 4.
tive 0; Mismatches
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                                                                                                                                                                       Version
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RESULT 57
US-08-758-306-141/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 35,353
REFERENCE/DOCKET NUMBER: 14
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 758-4800
TELEFAX: (212) 751-6849
INFORMATION FOR SEQ ID NO: 104:
                                INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
                                                                           ATTORNEY AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 212/132
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
                                                                                                                                                                                           CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE:
DESCRIPTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: JP 31
FILING DATE: 21-NOV-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                          SOFTWARE: FastSeq Version 1.5
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 1
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                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Stinchcomb, Dan T. APPLICANT: McSwiggen, James F
                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/758 FILING DATE: December 3, 1996
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET:
                                                             TELEFAX:
                                                                                                                                                                                FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                  90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                T: Suite 4700
Los Angeles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   41, Application US/08758306
5807743
                                                   67-3510
                                                                                                                                                                                                                                                                                                                                                                                                                                   California
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17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3: Lyon & Lyon
633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                 U.S.A.
                                                               (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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/desc = "synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GAMMA-CHAIN EXPRESSION
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Pred. No.
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                                                                                                          US-08-758-306-143
                                                                                                                                           TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 100.0%;
Matches 12; Conservative
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                                                                             Query Match
                                                   Matches
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COMPUTER READABLE FORM:
MEDIUM TYPE: 35" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                           FILING DATE: December 3, CLASSIFICATION: 514 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                               TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: FastSeq Version 1.5
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
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TOPOLOGY: li
                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 21:
                                                                                                                                                                                                                                                                            NAME: Warburg, Richa
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM CON
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: U.S.A.
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                                                                                                                                  STRANDEDNESS:
                                                                                                                                                                                                                    TELEFAX:
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                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
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                      222 GCTGAACAACAA 233
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                                                   12;
                                                                Similarity 100.0%;
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California
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633 West Fifth
                                                                                                                                                                                                                    (213) 955-0440
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McSwiggen, James A.
VENTION: METHOD AND REAGENT FOR THE
                                                   Conservative
                                                                                                                    linear
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                                                                                                                                                                                                                                                                                     Richard J
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                                                                Score 12; DB 1; 
; Pred. No. 1.6e+
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                                                                                                                                                                                                                                                              212/132
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Pred. No.
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                                                   Mismatches
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1.6e+04;
                                                                  1.6e+04;
                                                                             Length 17;
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RESULT 59
US-08-758-306-145/c
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US-08-758-306-411/c
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                                                                                                                                                                                                     Sequence 411, Application US/08758306 Patent No. 5807743
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OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FRASESEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/758,306
FILING DATE: December 3, 1996
CLASSIFICATION E 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
FILING DATE:
                                                                                                                                                                                    GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 212/132
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                  APPLICANT: Stinchcomb, Dan T.
APPLICANT: McSwiggen, James A.
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: TREATMENT OF DISEASES
TITLE OF INVENTION: INTERLEUKUN-2 RECEPTOR
TITLE OF INVENTION: GAMMA-CHAIN EXPRESSION
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
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ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: sing
TOPOLOGY: linear
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STREET: Suite 4700
CITY: Los Angeles
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ADDRESS:
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INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
LENGTH: nucleic acid
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequence 413, Application US/08758306
Patent No. 5807743
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 12; Conserv
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NAME: WAZDUTG, RICHARD J.
REGISTRATION UNMER: 32,327
REFERENCE/DOCKET NUMBER: 212/
TELECONMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US
FILING DATE: December
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS
SOFTWARE: FastSeq Version 1.5
CURRENT APPLICATION DATA:
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MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
SOFTWARE: FastSeq Version 1.5 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
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                                                   COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C
                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
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TOPOLOGY: lir
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633 West Fifth Street
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Stinchcomb, Dam.
Stinchcomb, Dames A.
The Authority Authori
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100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              METHOD AND REAGENT FOR TREATMENT OF DISEASES ASSOCIATED WITH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GAMMA-CHAIN EXPRESSION
                                                                     IBM P.C. DOS 5.0
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US-08-584-040-4138/c
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INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
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                                                                       APPLICATION NUMBER: US/08/584,0
FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: OCCODET 26, 1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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COMPUTER READABLE FORM:
MEDIUM TYPE: 35." Diskette, 1.44 Mb
MEDIUM TYPE: storage
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NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 212/132
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                   COMPUTER: 15".
OPERATING SYSTEM:
OPERATING SYSTEM:
OPERATING SYSTEM:
OPERATING SYSTEM:
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APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: TREATMENT OF DISEASES OR
TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                             CURRENT APPLICATION DATA:
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APPLICANT: McSwiggen, James
APPLICANT: Stinchcomb, Dan T.,
APPLICANT: Escobedo, Jaime
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APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                           STREET:
                                                                                                                                                                                                                                                                                                                                                                                     CITY: Los Angeles
STATE: California
                                       NAME: Warburg, Richard J. REGISTRATION NUMBER: 32,3
                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
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5. 6346398
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633 West Fifth Street
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                                                                                                                                                                     UMBER: US/08/584,040
January 11, 1996
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Perfect 5.1
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US-09-673-809-95/c
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LENGTH: 17
TYPE: DNA
ORGANISM: Homo sapiens
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
SOFTWARE: PatentIn version 3.0 SEQ ID NO 1905
                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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Patent No. 6566127
GENERAL ***
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Matches
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                                               TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions R. TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor FILE REFERENCE: MBHB00,876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR PILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1995-01-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Method for typing of FILE REFERENCE: PCT99.86.HLA. CURRENT APPLICATION NUMBER: US/09/673,809 CURRENT FILING DATE: 2000-10-20 PRIOR APPLICATION NUMBER: 98870088.6 PRIOR FILING DATE: 1998-04-20 NUMBER OF SEQ ID NOS: 107
                                                                                                                                                                                                                                                   APPLICANT:
                                       NUMBER OF SEQ ID NOS: 14225
                                                                                                                                                                                                                                                                                         APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Pavco, Pam
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Stinchcomb, Dan
Escobedo, Jaime
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thes 0;
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ORGANISM: Homo sapiens

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US-09-371-772B-5232/c
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; ORGANISM: Homo sapiens
US-09-371-772B-5231
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US-09-371-772B-5231/c
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; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-371-772B-5232
                                                                                       APPLICANT: Stinchoomb, Dan
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions I
TITLE OF INVENTION: Method and Reagent for the Treatment Factor Receptor
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION I
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APPLICANT: Escobedo, Jaime
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00,976-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-01-08
PRIOR FILING DATE: 1996-01-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
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APPLICANT: Pavco, Pam
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100.0%; Pred. No.
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US-09-371-772B-5233/c
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 6628
; LENGTH: 17
; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-371-772B-6628
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-371-772B-6628/c
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TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00,876-J (237/198)
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-01-08
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
NUMBER: OF SEQ ID NOS: 14225
SOFTWARE: PatentIn version 3.0
SEQ ID NO 5233
LENGTH: 17
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Best Local Similarity
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APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Retire OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00, 876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/05,974
PRIOR APPLICATION NUMBER: US 60/584,040
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-01-08
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APPLICANT:
APPLICANT:
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APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
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                                                                                                                                       NUMBER OF SEQ ID NOS: 14225
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ORGANISM: Homo sapiens
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Stinchcomb, Dan
Escobedo, Jaime
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100.0%; Pred. No. 1.6e+04
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RESULT 70
US-09-866-108A-7838/c
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US-09-854-140-12/c
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CURRENT APPLICATION NUMBER: US/09/854,140
CURRENT FILING DATE: 2001-05-11
NUMBER OF SEQ ID NOS: 14
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 12
LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity Matches 12; Conserv
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Best Local
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APPLICANT:
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Patent No. 666405
                                                                                                                                                                                                      FILE REFERENCE: ABOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
                                                                                                          PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR PRIOR PRIOR PRIOR APPLICATION NUMBER: PCT/US01/00667
                                                     PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
FILE REFERENCE: AEOMICA-7
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APPLICANT: JI, Yonggang
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                   APPLICATION NUMBER: PCT/US01/00669 FILING DATE: 2001-01-30
 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00665
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5. 6686188
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HANZEL, David K.
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                                                                                                                                                                                                                                                                                                                                                                             CHEN, Wensheng
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                                                                          ; ORGANISM: Homo US-09-866-108A-7844
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                                                                                                                                                 SOFTWARE: Aeomica
Patent No. 6686188
SEQ ID NO 7844
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NUMBER OF SEQ ID NOS: 15755
SOFTWARE: Acomica Sequence Listing Engine
Patent No. 6686188
SEQ ID NO 7838
LENGTH: 17
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PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
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CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
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APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, MAIK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
PRIOR.APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
                                                                                                           LENGTH: 17
TYPE: DNA
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PRIOR TILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
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                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: PCT/US01/00663 PRIOR FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00668
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FILING DATE: 2001-01-30
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Similarity 100.0%; F
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N, Sharron
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Pred. No.
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Patent No. 60002.
Patent No. 60002.
Patent No. 60002.
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SEQ ID NO 10378
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                                                                                                     APPLICANT:
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APPLICANT: JI, Yongga
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
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PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
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CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
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APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron
APPLICANT: HANZEL, David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 60/236,359
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
                                              APPLICANT:
                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Homo sapiens
PPLICANT: SHANNON, Mark
ITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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APPLICATION NUMBER: PCT/US01/00663
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FILING DATE: 2001-01-30
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                                                         PENN, Sharron G.
HANZEL, David K.
RANK, David R.
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                                      CHEN, Wensheng
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                                                                                                                Yonggang
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100.0%; Pred. No. 1.6e+(
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RESULT 74
US-09-866-108A-10380/c
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                                           CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-01-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00664
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SEQ ID NO 10379
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CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
DBIOD STITUTE NAME: 2000-10-04
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION UMMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
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PRIOR APPLICATION NUMBER: PCT/US01/00663
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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APPLICATION NUMBER: PCT/US01/00669
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APPLICATION NUMBER: PCT/US01/00665
APPLICATION NUMBER: PCT/US01/00669
FILING DATE: 2001-01-30
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Similarity 100.0%;
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RANK, David R.
CHEN, Wensheng
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Pred. No.
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APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN
FILE REPERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR APPLICATION NUMBER: PCT/US01/00669
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US-09-866-108A-10381/c
; Sequence 10381, Application US/09866108A
; Patent No. 6686188;
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; ORGANISM: Homo sapiens
US-09-866-108A-10380
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NUMBER OF SEQ ID NOS: 15755
SOFTWARE: Acomica Sequence Listing Engine
Patent No. 6686188
SEQ ID NO 10381
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Patent No. 6666188
SEQ ID NO 10380
LENGTH: 17
Query Match 2.4%; Score 12; DB 4; Length 17; Best Local Similarity 100.0%; Pred. No. 1.6e+04; Matches 12; Conservative 0; Mismatches 0; Indels
                                                                                                                                TYPE: DNA
ORGANISM: Homo mapienm
3-09-866-108A-10381
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PRIOR APPLICATION NUMBER: PCT/USO1/00663
PRIOR FILING DATE: 2001-01-30
Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 15755
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APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
APPLICANT: RANK, David R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
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